



SEQUENCE LISTING

RECEIVED
#3

AUG 08 2001

TECH CENTER 1600/2900

<110> Ruvkun, Gary
Kimura, Koutarou
Patterson, Garth
Ogg, Scott
Paradis, Suzanne
Tissenbaum, Heidi
Morris, Jason
Koweek, Allison

<120> THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
IMPAIRED GLUCOSE TOLERANCE CONDITIONS

<130> 00786/351005

<140> US 09/844,353
<141> 2001-04-27

<150> US 08/857,076
<151> 1997-05-15

<160> 114

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 1
cgctacggca aaaaagtgaa

20

<210> 2
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 2
cgatgatgaa gatacccc

18

<210> 3
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 3
tgatgcgaac ggcgatcgat 20

<210> 4
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 4
acgctggatc atctacatta 20

<210> 5
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 5
ggttaattt cccaaatgg ag 22

<210> 6
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 6
gctcacgggt cacacaacga 20

<210> 7
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 7
tgatgcgaac ggcgatcgat 20

<210> 8
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 8
tgaggggccaa ctaaagaaga c 21

10 20 30 40 50 60 70 80 90 100 110 120 130 140 150

<210> 9
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 9
cgctacggca aaaaagtgaa 20

<210> 10
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer/probe derived from C. elegans

<400> 10
gacgatcccg aggtgagtagt 20

<210> 11
<211> 5816
<212> DNA
<213> Caenorhabditis elegans

<220>
<221> misc_feature
<222> (1)...(5816)
<223> n = A,T,C or G

<400> 11
ggtttaatta cccaagtttg agctccaaga gcacacatct gatcgtcgga ttctactgta 60
ctccccgaaa aaccaacaaa aaacacaagt tttgaacac ttgtaaatgc agacagaacg 120
atgacgagaa tgaatatattgt cagatgtcg agacgacaca aaattttgaa aaattttgaa 180
gaagagaatc tcggcccgag ctgctcgctc acgacttcaa caaccgctgc caccgaagct 240
ctcggaaacaa ccactgagga tatgaggcctt aagcagcagc gaagctcgtc gcgtgcacag 300
gagcacgata ttgtcgacgg caatcaccac gacgacgagc acatcacaat gagacggctt 360
cgacttgtca aaaattcgcg gacgcggcgt agaacgcgc ccgattcaag tatggactgc 420
tatgagggaaa acccgccatc aaaaaaactt caataaaatta ttcttggatt tctaaaaagt 480
catcaatgac gtcattaaatg cttttactgc tattcgctt tgtacagccg tgcctcaa 540
tagtcgaaaa acgatgcggc ccaatcgata ttgcggat tagtgcggat attaaggccgc 600
aatggtcgaa acttgggtat ccgaacgaaa aagattggc tggtcagaga atggtaact 660
gcacagtgtt ggaagggttcg ctgacaatct catttgtact gaaacacaaag acaaaagcac 720
aagaagaaat gcatcgaatg ctacagccaa gatattccca agacgaattt atcactttc 780
cgcatctacg tgaattact ggaactctgc tcgttttgc gactgagga ttgtggatt 840
tgcgtaaaaat ttccccaat ctgcgtgtaa ttggaggccg ttgcgtgat caacactatg 900
cgctgataat ttatcgaaat ccggattttgg agatcggtct tgacaagctt tccgttaattc 960
gaaatgggtgg tgtacggata atcgataatc gaaaactgtg ctacacgaaa acgattgatt 1020
ggaaacattt gatcacttct tccatcaacg atgttgcgt tgataatgc gccgagtagc 1080
ctgtcactga gactggattt atgtgccac gtggagctt cgaagaggat aaaggcgaat 1140
caaagtgtca ttatttggag gaaaagaatc aggaacaagg tgcgttgc gttcagatgtt 1200
gttggtcgaa caccacttgc caaaagtctt gtgcgttgc tcgttgc tcaacgaaag 1260
aaatcgacc gggatgtat gcgacggcg atcgatgtca cgatcaatgc gtggccgtt 1320
gtgagcgtgt gaatgtatgcc acagcatgcc acgcgtgcaa gaatgtctat cacaaggaa 1380
agtgtatcga aaagtgtat gctcacctgtt accttctcc tcaacggtcg tgcgtgaccc 1440
gtgagcgtgt tctgcagctg aatccgggtgc tctcgaaacaa aacagtgcct atcaaggcga 1500

ପାଇଁ କିମ୍ବା କିମ୍ବା କିମ୍ବା କିମ୍ବା କିମ୍ବା କିମ୍ବା

cgccggccct ttgcgtcggat aaatgtcccg atggtatca aatcaaccg gatgatcatc 1560
gagaatgccg aaaatgcgtt ggcaagtgtg agattgtgtg cgagatcaat cacgtcattg 1620
atacgtttcc gaaggcacag gcgatcaggg tatgaatat tattgacgaa aatctgacga 1680
tcgagatcc cgaaaacag gattcgggaa tggcgtccgaa gttgaaggat atatttgcga 1740
acattcacac gatcaccggc tacctgttgg tacgtcaatc gtcaccgtt atctcggtga 1800
acatgttccg gaatttacga cgtattgagg caaagtcaact gttcagaaat ctatatgcta 1860
tcacagttt tgaaaatccg aattaaaaa agcttacga ttcaacgacg gatttgacgc 1920
ttgatcggtg aactgtgtca attgccaata acaagatgtt atgcttcaag tatatoaagc 1980
agctaatgtc aaagttaat ataccactcg atccgataga tcaatcagaa gggacaatg 2040
gtgagaaggn aatctgtgag gatatggcaa tcaacgttag catcacagcg gtcaacgcgg 2100
actcggtctt cttagtggg ccctcattca acattaccga tatagatcgat cgaardttc 2160
tcggctacga gctcttc taaagaagttc cacgaatcgat tgagaacatg acgatcgaag 2220
aggatcgaa tgcgtgtgtc gattcgtggc agagtgtctt caaacagtag tacgagacgt 2280
cgaacgggtga accgacccccg gacattttt tggatattgg accgcgcgag cgaattcggc 2340
cgaatacgct ctacgcgtac tatgtggcga cgcagatgtt gttgcgtc ggtgcgaaga 2400
acgggtgtatc gaagattggg ttgttgagga cgagactata tacgcgtat cctccgcgt 2460
tggcactagc gcaatcgat tggacgtca ttcatattac gtggaaagcg cgcgtcaac 2520
cgaacggaga cctcacgcatacataattt tggcgtgtga gaatgaagtg agcccgtagc 2580
aggaagccga aaagtttgtt acagatcgaa gcaccccccgc aatcgcacaa cgcacgaa 2640
atccgaaaga gacgattgtatcgatataattt tccgtcatca cgtaccgttag 2700
ctccgcacact ttgtactatg atgggtcraig aagatcagca gaaaacgtgc gtcgcaacgc 2760
ccgggttgg ttcgtgttcg gctatcgaa aatcatcgga acagaacaag aagaaggcgc 2820
cgatccgat gtcggcgtatc gaatcatctg catttgagaa taagctgtt gatgaggtt 2880
taatgcccgg agacacgcatacataattt gatcaattga agacgcgaat cgagtcaatg 2940
aagagttggaaaagctgaa aattttggaa aagctccaaa aactctcggt gaaaagaagc 3000
cgctgatcca tatttcgaa aagaagccgt cgagcagcag caccacatcc acaccggctc 3060
caacgatcgc atcaatgtat gccttaacaa gaaacccgcg tacgggtcccc ggaacaagga 3120
ttcggtcttca cgagatctac gaaaccttac cgggaagctg ggcgattaaat gtatcagctc 3180
tggcattggta taatagttat gtgatacggaa atttgaagca ttacacactt tatgcgattt 3240
ctctatccgc gtggccaaaac atgacagtagc cgggagcatc ttgctcaata tcccattcg 3300
cgggagcatt gaaacgaaca aaacacatca cagacattga taaaagtgtt aatgaaacaa 3360
ttgaatggag atttatgaat aatagtcaac aagtcaacgt gacgtgggat ccaccgactg 3420
aagtgaatgg tggaaatattt gtttatgtt taaagcttac gtcacaaagtc gatggatcaa 3480
ttgttatgac gagatgtgtc ggtgcgaaga gaggatattc aacacggaa cagggtgtcc 3540
tattccagaa tttggccgat ggacgttatt ttgtctcagt aacggcgcacc tctgtacacg 3600
gchgctggacc ggaagccgaa tcctccgacc caatcgtcgat catgacgcca ggcttctca 3660
ctgtggaaat catttcgcg atgcttctcg tcttttgc ttaatgtca attgcccgtt 3720
gtataatcta ctactacatt caagtacgct acggccaaaaa aagtggaaatct ctatctact 3780
ttatgcattt gaatcccgaa tattgtgtgg acaataagta caatgcacgac gattgggagc 3840
tacggcagga tgatgtgtg ctggacaaac agtgggaga gggatcattt gggaaatgtt 3900
accttaggaac tggaaataat gtttttctc tgatgggtga tcgtttccgaa ccgtgtgc 3960
ttaagatata tttgtatgtat ccagcgtcgat ctgagaatct caactatctc atggaaatcta 4020
atattatgaa gaactttaag actaacttta tcgtcacta gtaatgggat atctctactg 4080
tacaaccgcg gatgttgcg atggaaatgtt tgatcttgg aatctccgt gactatctcc 4140
gatcgaaacgc cgaagacgaa gtgttcaatg agacggactg caacttttc gacataatcc 4200
cgaggggataattccatcgat tggccgcac agattgtga tgatggcgttacccgtt 4260
cgctcaagtttgcgtcgat gatctcgccg cacgtatttgcatgataat cgggatgaga 4320
ctgtcaagat tggagatttc ggaatggctc gtgatcttattt ctatcatgac tattataagc 4380
catggccaa gcgatgtatc cctgttgcgtt gatgtcacc cgagtcgtt gaaagacggaa 4440
agtttgcactc gaaatctgtat gtttggagct tcggagttgt tctctatgaa atggttacac 4500
tcgggtgcata gccatataattt gtttggatgta atgatggat gttgaattt atggaaatgg 4560
cccgaaatgtt tataaagaag cccgaatgtt gtggaaacta ttggtataag gtatgaaaa 4620
tgtgtcgatc atactcacctt cgggatcgatc cgacgttctt ccagtcgtt catcttctag 4680
cagctgaagc ttccaccagaa ttccgagatt tatttttttgc tcttgcgtt gaaatctgtatc 4740
tccttgacga ttcaagaagca ctggatcttgcg atgatattgtt tgatgtatgat atgaatgtc 4800
aggttgcgtc ggtggcaccgc gatgttgcg acgtcgatgtt tcagactgtat tcggaaacgc 4860
ggaataacggc ttcaataccg ttgaaacagt ttaagacgtt ccctccgcattt aatgcgacga 4920
cgagtcattt gacaatatcg attgtatgaga caccgtgaa agcgaagcagc cgagaaggat 4980

EQUUS EQUUS

cgctggatga ggagtacgca ttgatgaatc atagtggagg tccgagtgat gcggaagttc 5040
ggacgtatgc tggatggaa gattatgtgg agagagatgt tcgagagaat gatgtgccaa 5100
cgcgacaaaa tactggtgc tcaacatcaa gttacacagg tgggtgtcca tattgcctaa 5160
caaatacgatgg tggatggaaat gaacgaggag ccgggttcgg tgaaggcgtcg cgttactg 5220
atggatggatgg aagtggacat ttaaatgtatgt tgaaaaagag atatcatcca 5280
tggatacgcg ccggagcaccg ggcgcctcga gctttccta cgggtgttcca cagacgaatt 5340
ggagtggaaa tcgtggtgc acgttattat cgagtaaagc tcaacaggca gcaactgcag 5400
cagcagcagc agcagcagct ctccaacagc aacaaaatgg tggtcgaggc gatcgattaa 5460
ctcaactacc cggaaactgga catttacaat cgacacgtgg tggacaagat ggagattata 5520
ttgaaactga accaaaaat tatagaaata atggatctcc atcgcgaaac ggcaacagcc 5580
gtgacattt caacggacgt tcggcttcg gtgaaaatga gcatctaatic gaggataatg 5640
agcatcatcc acttgtctga aacccccaaa aaatccgc tcttaaatta taaattatct 5700
cccacattat catactctca cacgaatatac ggatttttt tctgaaaaat 5760
tctgaataat tttacccat tttcaaaatc tctgtatccc ttttggat taccgg 5816

<210> 12
<211> 1724
<212> PRT
<213> Caenorhabditis elegans

<400> 12
Met Thr Ser Leu Met Leu Leu Leu Leu Phe Ala Phe Val Gln Pro Cys
1 5 10 15
Ala Ser Ile Val Glu Lys Arg Cys Gly Pro Ile Asp Ile Arg Asn Arg
20 25 30
Pro Trp Asp Ile Lys Pro Gln Trp Ser Lys Leu Gly Asp Pro Asn Glu
35 40 45
Lys Asp Leu Ala Gly Gln Arg Met Val Asn Cys Thr Val Val Glu Gly
50 55 60
Ser Leu Thr Ile Ser Phe Val Leu Lys His Lys Thr Lys Ala Gln Glu
65 70 75 80
Glu Met His Arg Ser Leu Gln Pro Arg Tyr Ser Gln Asp Glu Phe Ile
85 90 95
Thr Phe Pro His Leu Arg Glu Ile Thr Gly Thr Leu Leu Val Phe Glu
100 105 110
Thr Glu Gly Leu Val Asp Leu Arg Lys Ile Phe Pro Asn Leu Arg Val
115 120 125
Ile Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg
130 135 140
Asn Pro Asp Leu Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn
145 150 155 160
Gly Gly Val Arg Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr
165 170 175
Ile Asp Trp Lys His Leu Ile Thr Ser Ser Ile Asn Asp Val Val Val
180 185 190
Asp Asn Ala Ala Glu Tyr Ala Val Thr Glu Thr Gly Leu Met Cys Pro
195 200 205
Arg Gly Ala Cys Glu Glu Asp Lys Gly Glu Ser Lys Cys His Tyr Leu
210 215 220
Glu Glu Lys Asn Gln Glu Gln Gly Val Glu Arg Val Gln Ser Cys Trp
225 230 235 240
Ser Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro
245 250 255
Thr Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys His
260 265 270
Asp Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala Cys

EQUINE

275	280	285
His Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys Cys		
290	295	300
Asp Ala His Leu Tyr Leu Leu Gln Arg Arg Cys Val Thr Arg Glu		
305	310	315
Gln Cys Leu Gln Leu Asn Pro Val Leu Ser Asn Lys Thr Val Pro Ile		
325	330	335
Lys Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr Gln		
340	345	350
Ile Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys Cys		
355	360	365
Glu Ile Val Cys Glu Ile Asn His Val Ile Asp Thr Phe Pro Lys Ala		
370	375	380
Gln Ala Ile Arg Leu Cys Asn Ile Ile Asp Gly Asn Leu Thr Ile Glu		
385	390	395
Ile Arg Gly Lys Gln Asp Ser Gly Met Ala Ser Glu Leu Lys Asp Ile		
405	410	415
Phe Ala Asn Ile His Thr Ile Thr Gly Tyr Leu Leu Val Arg Gln Ser		
420	425	430
Ser Pro Phe Ile Ser Leu Asn Met Phe Arg Asn Leu Arg Arg Ile Glu		
435	440	445
Ala Lys Ser Leu Phe Arg Asn Leu Tyr Ala Ile Thr Val Phe Glu Asn		
450	455	460
Pro Asn Leu Lys Lys Leu Phe Asp Ser Thr Thr Asp Leu Thr Leu Asp		
465	470	475
Arg Gly Thr Val Ser Ile Ala Asn Asn Lys Met Leu Cys Phe Lys Tyr		
485	490	495
Ile Lys Gln Leu Met Ser Lys Leu Asn Ile Pro Leu Asp Pro Ile Asp		
500	505	510
Gln Ser Glu Gly Thr Asn Gly Glu Lys Ala Ile Cys Glu Asp Met Ala		
515	520	525
Ile Asn Val Ser Ile Thr Ala Val Asn Ala Asp Ser Val Phe Phe Ser		
530	535	540
Trp Pro Ser Phe Asn Ile Thr Asp Ile Asp Gln Arg Lys Phe Leu Gly		
545	550	555
Tyr Glu Leu Phe Phe Lys Glu Val Pro Arg Ile Asp Glu Asn Met Thr		
565	570	575
Ile Glu Glu Asp Arg Ser Ala Cys Val Asp Ser Trp Gln Ser Val Phe		
580	585	590
Lys Gln Tyr Tyr Glu Thr Ser Asn Gly Glu Pro Thr Pro Asp Ile Phe		
595	600	605
Met Asp Ile Gly Pro Arg Glu Arg Ile Arg Pro Asn Thr Leu Tyr Ala		
610	615	620
Tyr Tyr Val Ala Thr Gln Met Val Leu His Ala Gly Ala Lys Asn Gly		
625	630	635
Val Ser Lys Ile Gly Phe Val Arg Thr Ser Tyr Tyr Thr Pro Asp Pro		
645	650	655
Pro Thr Leu Ala Leu Ala Gln Val Asp Ser Asp Ala Ile His Ile Thr		
660	665	670
Trp Glu Ala Pro Leu Gln Pro Asn Gly Asp Leu Thr His Tyr Thr Ile		
675	680	685
Met Trp Arg Glu Asn Glu Val Ser Pro Tyr Glu Glu Ala Glu Lys Phe		
690	695	700
Cys Thr Asp Ala Ser Thr Pro Ala Asn Arg Gln Arg Thr Lys Asp Pro		
705	710	715
Lys Glu Thr Ile Val Ala Asp Lys Pro Val Asp Ile Pro Ser Ser Arg		
725	730	735
Thr Val Ala Pro Thr Leu Leu Thr Met Met Gly His Glu Asp Gln Gln		

H G D E F S T C P Q R N V I M

740	745	750
Lys Thr Cys Ala Ala Thr Pro Gly Cys Cys Ser Cys Ser Ala Ile Glu		
755	760	765
Glu Ser Ser Glu Gln Asn Lys Lys Lys Arg Pro Asp Pro Met Ser Ala		
770	775	780
Ile Glu Ser Ser Ala Phe Glu Asn Lys Leu Leu Asp Glu Val Leu Met		
785	790	795
Pro Arg Asp Thr Met Arg Val Arg Arg Ser Ile Glu Asp Ala Asn Arg		
805	810	815
Val Ser Glu Glu Leu Glu Lys Ala Glu Asn Leu Gly Lys Ala Pro Lys		
820	825	830
Thr Leu Gly Gly Lys Lys Pro Leu Ile His Ile Ser Lys Lys Lys Pro		
835	840	845
Ser Ser Ser Ser Thr Thr Ser Thr Pro Ala Pro Thr Ile Ala Ser Met		
850	855	860
Tyr Ala Leu Thr Arg Lys Pro Thr Thr Val Pro Gly Thr Arg Ile Arg		
865	870	875
Leu Tyr Glu Ile Tyr Glu Pro Leu Pro Gly Ser Trp Ala Ile Asn Val		
885	890	895
Ser Ala Leu Ala Leu Asp Asn Ser Tyr Val Ile Arg Asn Leu Lys His		
900	905	910
Tyr Thr Leu Tyr Ala Ile Ser Leu Ser Ala Cys Gln Asn Met Thr Val		
915	920	925
Pro Gly Ala Ser Cys Ser Ile Ser His Arg Ala Gly Ala Leu Lys Arg		
930	935	940
Thr Lys His Ile Thr Asp Ile Asp Lys Val Leu Asn Glu Thr Ile Glu		
945	950	955
Trp Arg Phe Met Asn Asn Ser Gln Gln Val Asn Val Thr Trp Asp Pro		
965	970	975
Pro Thr Glu Val Asn Gly Gly Ile Phe Gly Tyr Val Val Lys Leu Lys		
980	985	990
Ser Lys Val Asp Gly Ser Ile Val Met Thr Arg Cys Val Gly Ala Lys		
995	1000	1005
Arg Gly Tyr Ser Thr Arg Asn Gln Gly Val Leu Phe Gln Asn Leu Ala		
1010	1015	1020
Asp Gly Arg Tyr Phe Val Ser Val Thr Ala Thr Ser Val His Gly Ala		
1025	1030	1035
Gly Pro Glu Ala Glu Ser Ser Asp Pro Ile Val Val Met Thr Pro Gly		
1045	1050	1055
Phe Phe Thr Val Glu Ile Ile Leu Gly Met Leu Leu Val Phe Leu Ile		
1060	1065	1070
Leu Met Ser Ile Ala Gly Cys Ile Ile Tyr Tyr Tyr Ile Gln Val Arg		
1075	1080	1085
Tyr Gly Lys Lys Val Lys Ala Leu Ser Asp Phe Met Gln Leu Asn Pro		
1090	1095	1100
Glu Tyr Cys Val Asp Asn Lys Tyr Asn Ala Asp Asp Trp Glu Leu Arg		
1105	1110	1115
Gln Asp Asp Val Val Leu Gly Gln Gln Cys Gly Glu Gly Ser Phe Gly		
1125	1130	1135
Lys Val Tyr Leu Gly Thr Gly Asn Asn Val Val Ser Leu Met Gly Asp		
1140	1145	1150
Arg Phe Gly Pro Cys Ala Ile Lys Ile Asn Val Asp Asp Pro Ala Ser		
1155	1160	1165
Thr Glu Asn Leu Asn Tyr Leu Met Glu Ala Asn Ile Met Lys Asn Phe		
1170	1175	1180
Lys Thr Asn Phe Ile Val Gln Leu Tyr Gly Val Ile Ser Thr Val Gln		
1185	1190	1195
Pro Ala Met Val Val Met Glu Met Met Asp Leu Gly Asn Leu Arg Asp		

卷之三

1205	1210	1215
Tyr Leu Arg Ser Lys Arg Glu Asp	Glu Val Phe Asn Glu Thr Asp Cys	
1220	1225	1230
Asn Phe Phe Asp Ile Ile Pro Arg Asp Lys Phe His Glu Trp Ala Ala		
1235	1240	1245
Gln Ile Cys Asp Gly Met Ala Tyr Leu Glu Ser Leu Lys Phe Cys His		
1250	1255	1260
Arg Asp Leu Ala Ala Arg Asn Cys Met Ile Asn Arg Asp Glu Thr Val		
1265	1270	1275
Lys Ile Gly Asp Phe Gly Met Ala Arg Asp Leu Phe Tyr His Asp Tyr		
1285	1290	1295
Tyr Lys Pro Ser Gly Lys Arg Met Met Pro Val Arg Trp Met Ser Pro		
1300	1305	1310
Glu Ser Leu Lys Asp Gly Lys Phe Asp Ser Lys Ser Asp Val Trp Ser		
1315	1320	1325
Phe Gly Val Val Leu Tyr Glu Met Val Thr Leu Gly Ala Gln Pro Tyr		
1330	1335	1340
Ile Gly Leu Ser Asn Asp Glu Val Leu Asn Tyr Ile Gly Met Ala Arg		
1345	1350	1355
Lys Val Ile Lys Lys Pro Glu Cys Cys Glu Asn Tyr Trp Tyr Lys Val		
1365	1370	1375
Met Lys Met Cys Trp Arg Tyr Ser Pro Arg Asp Arg Pro Thr Phe Leu		
1380	1385	1390
Gln Leu Val His Leu Leu Ala Ala Glu Ala Ser Pro Glu Phe Arg Asp		
1395	1400	1405
Leu Ser Phe Val Leu Thr Asp Asn Gln Met Ile Leu Asp Asp Ser Glu		
1410	1415	1420
Ala Leu Asp Leu Asp Asp Ile Asp Asp Thr Asp Met Asn Asp Gln Val		
1425	1430	1435
Val Glu Val Ala Pro Asp Val Glu Asn Val Glu Val Gln Ser Asp Ser		
1445	1450	1455
Glu Arg Arg Asn Thr Asp Ser Ile Pro Leu Lys Gln Phe Lys Thr Ile		
1460	1465	1470
Pro Pro Ile Asn Ala Thr Thr Ser His Ser Thr Ile Ser Ile Asp Glu		
1475	1480	1485
Thr Pro Met Lys Ala Lys Gln Arg Glu Gly Ser Leu Asp Glu Glu Tyr		
1490	1495	1500
Ala Leu Met Asn His Ser Gly Gly Pro Ser Asp Ala Glu Val Arg Thr		
1505	1510	1515
Tyr Ala Gly Asp Gly Asp Tyr Val Glu Arg Asp Val Arg Glu Asn Asp		
1525	1530	1535
Val Pro Thr Arg Arg Asn Thr Gly Ala Ser Thr Ser Ser Tyr Thr Gly		
1540	1545	1550
Gly Gly Pro Tyr Cys Leu Thr Asn Arg Gly Gly Ser Asn Glu Arg Gly		
1555	1560	1565
Ala Gly Phe Gly Glu Ala Val Arg Leu Thr Asp Gly Val Gly Ser Gly		
1570	1575	1580
His Leu Asn Asp Asp Asp Tyr Val Glu Lys Glu Ile Ser Ser Met Asp		
1585	1590	1595
Thr Arg Arg Ser Thr Gly Ala Ser Ser Ser Tyr Gly Val Pro Gln		
1605	1610	1615
Thr Asn Trp Ser Gly Asn Arg Gly Ala Thr Tyr Tyr Thr Ser Lys Ala		
1620	1625	1630
Gln Gln Ala Ala Thr Ala Ala Ala Ala Ala Ala Leu Gln Gln		
1635	1640	1645
Gln Gln Asn Gly Gly Arg Gly Asp Arg Leu Thr Gln Leu Pro Gly Thr		
1650	1655	1660
Gly His Leu Gln Ser Thr Arg Gly Gly Gln Asp Gly Asp Tyr Ile Glu		

1665 1670 1675 1680
Thr Glu Pro Lys Asn Tyr Arg Asn Asn Gly Ser Pro Ser Arg Asn Gly
1685 1690 1695
Asn Ser Arg Asp Ile Phe Asn Gly Arg Ser Ala Phe Gly Glu Asn Glu
1700 1705 1710
His Leu Ile Glu Asp Asn Glu His His Pro Leu Val
1715 1720

<210> 13
<211> 139
<212> PRT
<213> Caenorhabditis elegans

<400> 13
Thr Ser Gly Ser Gly Met Gly Pro Thr Thr Leu His Lys Leu Thr Ile
1 5 10 15
Gly Gly Gln Ile Arg Leu Thr Gly Arg Val Gly Ser Gly Arg Phe Gly
20 25 30
Asn Val Ser Arg Gly Asp Tyr Arg Gly Glu Ala Val Ala Val Lys Val
35 40 45
Phe Asn Ala Leu Asp Glu Pro Ala Phe His Lys Glu Thr Glu Ile Phe
50 55 60
Glu Thr Arg Met Leu Arg His Pro Asn Val Leu Arg Tyr Ile Gly Ser
65 70 75 80
Asp Arg Val Asp Thr Gly Phe Val Thr Glu Leu Trp Leu Val Thr Glu
85 90 95
Tyr His Pro Ser Gly Ser Leu His Asp Phe Leu Leu Glu Asn Thr Val
100 105 110
Asn Ile Glu Thr Tyr Tyr Asn Leu Met Arg Ser Thr Ala Ser Gly Leu
115 120 125
Ala Phe Leu His Asn Gln Ile Gly Gly Ser Lys
130 135

<210> 14
<211> 62
<212> PRT
<213> Caenorhabditis elegans

<400> 14
Glu Asp Ala Ala Ser Asp Ile Ile Ala Asn Glu Asn Tyr Lys Cys Gly
1 5 10 15
Thr Val Arg Tyr Leu Ala Pro Glu Ile Leu Asn Ser Thr Met Gln Phe
20 25 30
Thr Val Phe Glu Ser Tyr Gln Cys Ala Asp Val Tyr Ser Phe Ser Leu
35 40 45
Val Met Trp Glu Thr Leu Cys Arg Cys Glu Asp Gly Asp Val
50 55 60

<210> 15
<211> 31
<212> PRT
<213> Caenorhabditis elegans

<400> 15
Lys Pro Ala Met Ala His Arg Asp Ile Lys Ser Lys Asn Ile Met Val

1 5 10 15
Lys Asn Asp Leu Thr Cys Ala Ile Gly Asp Leu Gly Leu Ser Leu
20 25 30

<210> 16
<211> 72
<212> PRT
<213> Caenorhabditis elegans

<400> 16
Ile Pro Tyr Ile Glu Trp Thr Asp Arg Asp Pro Gln Asp Ala Gln Met
1 5 10 15
Phe Asp Val Val Cys Thr Arg Arg Leu Arg Pro Thr Glu Asn Pro Leu
20 25 30
Trp Lys Asp His Pro Glu Met Lys His Ile Met Glu Ile Ile Lys Thr
35 40 45
Cys Trp Asn Gly Asn Pro Ser Ala Arg Phe Thr Ser Tyr Ile Cys Arg
50 55 60
Lys Arg Met Asp Glu Arg Gln Gln
65 70

<210> 17
<211> 150
<212> PRT
<213> Caenorhabditis elegans

<400> 17
Tyr Phe Glu Ser Val Asp Arg Phe Leu Tyr Ser Cys Val Gly Tyr Ser
1 5 10 15
Val Ala Thr Tyr Ile Met Gly Ile Lys Asp Arg His Ser Asp Asn Leu
20 25 30
Met Leu Thr Glu Asp Gly Lys Tyr Val His Ile Asp Phe Gly His Ile
35 40 45
Leu Gly His Gly Lys Thr Lys Leu Gly Ile Gln Arg Asp Arg Gln Pro
50 55 60
Phe Ile Leu Thr Glu His Phe Met Thr Val Ile Arg Ser Gly Lys Ser
65 70 75 80
Val Asp Gly Asn Ser His Glu Leu Gln Lys Phe Lys Thr Leu Cys Val
85 90 95
Glu Ala Tyr Glu Val Met Trp Asn Asn Arg Asp Leu Phe Val Ser Leu
100 105 110
Phe Thr Leu Met Leu Gly Met Glu Leu Pro Glu Leu Ser Thr Lys Ala
115 120 125
Asp Leu Asp His Leu Lys Lys Thr Leu Phe Cys Asn Gly Glu Ser Lys
130 135 140
Glu Glu Ala Arg Lys Phe
145 150

<210> 18
<211> 113
<212> PRT
<213> Caenorhabditis elegans

<400> 18
Ser Pro Leu Asp Pro Val Tyr Lys Leu Gly Glu Met Ile Ile Asp Lys

1 5 10 15
Ala Ile Val Leu Gly Ser Ala Lys Arg Pro Leu Met Leu His Trp Lys
20 25 30
Asn Lys Asn Pro Lys Ser Asp Leu His Leu Pro Phe Cys Ala Met Ile
35 40 45
Phe Lys Asn Gly Asp Asp Leu Arg Gln Asp Met Leu Val Leu Gln Val
50 55 60
Leu Glu Val Met Asp Asn Ile Trp Lys Ala Ala Asn Ile Asp Cys Cys
65 70 75 80
Leu Asn Pro Tyr Ala Val Leu Pro Met Gly Glu Met Ile Gly Ile Ile
85 90 95
Glu Val Val Pro Asn Cys Lys Thr Ile Phe Glu Ile Gln Val Gly Thr
100 105 110
Gly

<210> 19

<211> 106

<212> PRT

<213> Caenorhabditis elegans

<400> 19

Leu Ala Phe Val Trp Thr Asp Arg Glu Asn Phe Ser Glu Leu Tyr Val
1 5 10 15
Met Leu Glu Lys Trp Lys Pro Pro Ser Val Ala Ala Leu Thr Leu
20 25 30
Leu Gly Lys Arg Cys Thr Asp Arg Val Ile Arg Lys Phe Ala Val Glu
35 40 45
Lys Leu Asn Glu Gln Leu Ser Pro Val Thr Phe His Leu Phe Ile Leu
50 55 60
Pro Leu Ile Gln Ala Leu Lys Tyr Glu Pro Arg Ala Gln Ser Glu Val
65 70 75 80
Gly Met Met Leu Leu Thr Arg Ala Leu Cys Asp Tyr Arg Ile Gly His
85 90 95
Arg Leu Phe Trp Leu Leu Arg Ala Glu Ile
100 105

<210> 20

<211> 139

<212> PRT

<213> Caenorhabditis elegans

<400> 20

Glu Tyr Trp Ile Val Thr Glu Phe His Glu Arg Leu Ser Leu Tyr Glu
1 5 10 15
Leu Leu Lys Asn Asn Val Ile Ser Ile Thr Ser Ala Asn Arg Ile Ile
20 25 30
Met Ser Met Ile Asp Gly Leu Gln Phe Leu His Asp Asp Arg Pro Tyr
35 40 45
Phe Phe Gly His Pro Lys Lys Pro Ile Ile His Arg Asp Ile Lys Ser
50 55 60
Lys Asn Ile Leu Val Lys Ser Asp Met Thr Thr Cys Ile Ala Asp Phe
65 70 75 80
Gly Leu Ala Arg Ile Tyr Ser Tyr Asp Ile Glu Gln Ser Asp Leu Leu
85 90 95
Gly Gln Val Gly Thr Lys Arg Tyr Met Ser Pro Glu Met Leu Glu Gly

Y F S C D E G Q H P K R N T S V W

	100		105		110										
Ala	Thr	Glu	Phe	Thr	Pro	Thr	Ala	Phe	Lys	Ala	Met	Asp	Val	Tyr	Ser
		115				120					125				
Met	Gly	Leu	Val	Met	Trp	Glu	Val	Ile	Ser	Arg					
		130				135									

<210> 21
<211> 61
<212> PRT
<213> Caenorhabditis elegans

<400> 21
Ile Gly Phe Asp Pro Thr Ile Gly Arg Met Arg Asn Tyr Val Val Ser
1 5 10 15
Lys Lys Glu Arg Pro Gln Trp Arg Asp Glu Ile Ile Lys His Glu Tyr
20 25 30
Met Ser Leu Leu Lys Lys Val Thr Glu Glu Met Trp Asp Pro Glu Ala
35 40 45
Cys Ala Arg Ile Thr Ala Gly Cys Ala Phe Ala Arg Val
50 55 60

<210> 22
<211> 20
<212> PRT
<213> Caenorhabditis elegans

<400> 22
Pro Ile Thr Asp Phe Gln Leu Ile Ser Lys Gly Arg Phe Gly Lys Val
1 5 10 15
Phe Lys Ala Gln
20

<210> 23
<211> 163
<212> PRT
<213> Caenorhabditis elegans

<400> 23
Thr Asp Ser Glu Thr Arg Ser Arg Phe Ser Leu Gly Trp Tyr Asn Asn
1 5 10 15
Pro Asn Arg Ser Pro Gln Thr Ala Glu Val Arg Gly Leu Ile Gly Lys
20 25 30
Gly Val Arg Phe Tyr Leu Leu Ala Gly Glu Val Tyr Val Glu Asn Leu
35 40 45
Cys Asn Ile Pro Val Phe Val Gln Ser Ile Gly Ala Asn Met Lys Asn
50 55 60
Gly Phe Gln Leu Asn Thr Val Ser Lys Leu Pro Pro Thr Gly Thr Met
65 70 75 80
Lys Val Phe Asp Met Arg Leu Phe Ser Lys Gln Leu Arg Thr Ala Ala
85 90 95
Glu Lys Thr Tyr Gln Asp Val Tyr Cys Leu Ser Arg Met Cys Thr Val
100 105 110
Arg Val Ser Phe Cys Lys Gly Trp Gly Glu His Tyr Arg Arg Ser Thr
115 120 125
Val Leu Arg Ser Pro Val Trp Phe Gln Ala His Leu Asn Asn Pro Met

130 135 140
His Trp Val Asp Ser Val Leu Thr Cys Met Gly Ala Pro Pro Arg Ile
145 150 155 160
Cys Ser Ser

<210> 24
<211> 44
<212> PRT
<213> Caenorhabditis elegans

<400> 24
Arg Ala Phe Arg Phe Pro Val Ile Arg Tyr Glu Ser Gln Val Lys Ser
1 5 10 15
Ile Leu Thr Cys Arg His Ala Phe Asn Ser His Ser Arg Asn Val Cys
20 25 30
Leu Asn Pro Tyr His Tyr Arg Trp Val Glu Leu Pro
35 40

<210> 25
<211> 38
<212> PRT
<213> Caenorhabditis elegans

<400> 25
Val Glu Tyr Glu Glu Ser Pro Ser Trp Leu Lys Leu Ile Tyr Tyr Glu
1 5 10 15
Glu Gly Thr Met Ile Gly Glu Lys Ala Asp Val Glu Gly His His Cys
20 25 30
Leu Ile Asp Gly Phe Thr
35

<210> 26
<211> 60
<212> PRT
<213> Caenorhabditis elegans

<400> 26
Asn Leu Ala Glu Thr Gly His Ser Lys Ile Met Arg Ala Ala His Lys
1 5 10 15
Val Ser Asn Pro Glu Ile Gly Tyr Cys Cys His Pro Thr Glu Tyr Asp
20 25 30
Tyr Ile Lys Leu Ile Tyr Val Asn Arg Asp Gly Arg Val Ser Ile Ala
35 40 45
Asn Val Asn Gly Met Ile Ala Lys Lys Cys Gly Cys
50 55 60

<210> 27
<211> 20
<212> PRT
<213> Caenorhabditis elegans

<400> 27
Asp Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala Tyr Met Cys Arg Gly

SEARCHED
SEARCHED
SEARCHED
SEARCHED
SEARCHED
SEARCHED
SEARCHED
SEARCHED
SEARCHED
SEARCHED

1	5	10	15
Asp Cys His Tyr			
20			

<210> 28
<211> 43
<212> PRT
<213> *Caenorhabditis elegans*

<400> 28
Val Cys Asn Ala Glu Ala Gln Ser Lys Gly Cys Cys Leu Tyr Asp Leu
1 5 10 15
Glu Ile Glu Phe Glu Lys Ile Gly Trp Asp Trp Ile Val Ala Pro Pro
20 25 30
Arg Tyr Asn Ala Tyr Met Cys Arg Gly Asp Cys
35 40

<210> 29
<211> 70
<212> PRT
<213> *Caenorhabditis elegans*

<400> 29
Asp Cys His Tyr Asn Ala His His Phe Asn Leu Ala Glu Thr Gly His
1 5 10 15
Ser Lys Ile Met Arg Ala Ala His Lys Val Ser Asn Pro Glu Ile Gly
20 25 30
Tyr Cys Cys His Pro Thr Glu Tyr Asp Tyr Ile Lys Leu Ile Tyr Val
35 40 45
Asn Arg Asp Gly Arg Val Ser Ile Ala Asn Val Asn Gly Met Ile Ala
50 55 60
Lys Lys Cys Gly Cys Ser
65 70

<210> 30
<211> 35
<212> PRT
<213> *Caenorhabditis elegans*

<400> 30
Cys Cys Leu Tyr Asp Leu Glu Ile Glu Phe Glu Lys Ile Gly Trp Asp
1 5 10 15
Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala Tyr Met Cys Arg Gly Asp
20 25 30
Cys His Tyr
35

<210> 31
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Degenerate probe

PDB ID: 1B3D

```
<221> misc_feature
<222> (1)...(23)
<223> n = A,T,C or G

<400> 31
ggntggayt rnrtnrtncc ncc 23

<210> 32
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Degenerate probe

<221> misc_feature
<222> (1)...(18)
<223> n = A,T,C or G

<400> 32
tgytgynnnnc cnacngar 18

<210> 33
<211> 127
<212> PRT
<213> Caenorhabditis elegans

<400> 33
Lys Phe His Glu Trp Ala Ala Gln Ile Cys Asp Gly Met Ala Tyr Leu
 1           5          10          15
Glu Ser Leu Lys Phe Cys His Arg Asp Leu Ala Ala Arg Asn Cys Met
 20          25          30
Ile Asn Arg Asp Glu Thr Val Lys Ile Gly Asp Phe Gly Met Ala Arg
 35          40          45
Asp Leu Phe Tyr His Asp Tyr Tyr Lys Pro Ser Gly Lys Arg Met Met
 50          55          60
Pro Val Arg Trp Met Ser Pro Glu Ser Leu Lys Asp Gly Lys Phe Asp
 65          70          75          80
Ser Lys Ser Asp Val Trp Ser Phe Gly Val Val Leu Tyr Glu Met Val
 85          90          95
Thr Leu Gly Ala Gln Pro Tyr Ile Gly Leu Ser Asn Asp Glu Val Leu
100         105         110
Asn Tyr Ile Gly Met Ala Arg Lys Val Ile Lys Lys Pro Glu Cys
115         120         125

<210> 34
<211> 131
<212> PRT
<213> Caenorhabditis elegans

<400> 34
Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro Thr
 1           5          10          15
Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys His Asp
 20          25          30
Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala Cys His
 35          40          45
```

Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys Cys Asp
50 55 60
Ala His Leu Tyr Leu Leu Gln Arg Arg Cys Val Thr Arg Glu Gln
65 70 75 80
Cys Leu Gln Leu Asn Pro Val Leu Ser Asn Lys Thr Val Pro Ile Lys
85 90 95
Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr Gln Ile
100 105 110
Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys Cys Glu
115 120 125
Ile Val Cys
130

<210> 35
<211> 103
<212> PRT
<213> Caenorhabditis elegans

<400> 35
Phe Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Leu Lys Asp Lys
1 5 10 15
Lys Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr
20 25 30
Lys Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu
35 40 45
Gln Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu
50 55 60
Trp Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His
65 70 75 80
Cys Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro
85 90 95
Tyr His Tyr Glu Ile Val Ile
100

<210> 36
<211> 79
<212> PRT
<213> Caenorhabditis elegans

<400> 36
Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val
1 5 10 15
Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr
20 25 30
Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val
35 40 45
Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys
50 55 60
Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe
65 70 75

<210> 37
<211> 106
<212> PRT
<213> Caenorhabditis elegans

BIOASSAY

<400> 37

Lys Lys Thr Thr Thr Arg Arg Asn Ala Trp Gly Asn Met Ser Tyr Ala
1 5 10 15
Glu Leu Ile Thr Thr Ala Ile Met Ala Ser Pro Glu Lys Arg Leu Thr
20 25 30
Leu Ala Gln Val Tyr Glu Trp Met Val Gln Asn Val Pro Tyr Phe Arg
35 40 45
Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly Trp Lys Asn Ser Ile Arg
50 55 60
His Asn Leu Ser Leu His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly
65 70 75 80
Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly
85 90 95
Met Asn Pro Arg Arg Thr Arg Glu Arg Ser
100 105

<210> 38

<211> 60

<212> PRT

<213> Caenorhabditis elegans

<400> 38

Glu Ile Lys Leu Ser Asp Phe Lys His Gln Leu Phe Glu Leu Ile Ala
1 5 10 15
Pro Met Lys Trp Gly Thr Tyr Ser Val Lys Pro Gln Asp Tyr Val Phe
20 25 30
Arg Gln Leu Asn Asn Phe Gly Glu Ile Glu Val Ile Phe Asn Asp Asp
35 40 45
Gln Pro Leu Ser Lys Leu Glu Leu His Gly Thr Phe
50 55 60

<210> 39

<211> 2784

<212> DNA

<213> Caenorhabditis elegans

<400> 39

atgaagctaa tagcaacttc tcttctagtt cccgacgagc acacaccgat gatgtcacca 60
gtgaatacaa ctacaaagat tctacaacgg agtggattta aatggaaat cccgcataat 120
ttggatccag acagtcagga ttagtaccgg gaagatgggt tcaactaccc ggatccagat 180
ttatttgaca caaaaaacac aaatatgacc gagtacgatt tggatgtgtt gaagcttgg 240
aaaccagcag tagatgaagc acggaaaaag atcgaagtcc ccgacgctag tgcgcggcca 300
aacaaaattt tagaatattt gatgtattat agaacgtta aagaaagtga actcatacaa 360
ctgaatgcgt atcgacaaa acgaaatcga ttatcgttga acttggtcaa aaacaatatt 420
gatcgagagt tcgacaaa agcttgcgag tcccttgcgtt aaaaatttggaa ggataagaag 480
aatgatctcc agaacctgtat tgatgtgtt ctttcaaaag gtacaaaat taccgggttgc 540
attacaattt caaggacact tgatggccgg ttacagggtcc acggaaagaaaa aggtttccct 600
cacgttgtt atggcaact gtggagggtt aatgaaatcga caaaaaacgaa aacgcgtcat 660
gtggaccact gcaagcacgc atttggaaatg aaaagtgaca tggtatgcgt gaatccctat 720
caactacgaaa ttgtcattgg aactatgatt gttggcgcaga gggatcatga caatcgagat 780
atgccgcgc cacatcaacg ctaccacact ccagggtccgc aggatccagt tgacgatatg 840
agtagattt taccaccaggc ttccattcgt ccgcctccga tgaacatgcaca aacaaggcct 900
cagcctatgc ctcaacaatt gccttcagg ggcgcacatg ttgcccattcc tctccacat 960
caggcgccac ataacccagg ggtttcacat ccgtactcca ttgctccaca gaccattac 1020
ccgttgaaca tgaacccaat tccgcaaattt ccgcacaaatgc cacaatgcc accacatctc 1080
catcaggat atgaaatgaa tggccgagtt tgcttccatg aaaacaacaa tccattccac 1140

caaaaatcacc attataatga tattagccat ccaaatactact attcctacga ctgtggtccg 1200
 aacttgtacg ggtttccaac tccttatccg gattttcacc atcctttcaa tcagcaacca 1260
 caccagccgc cacaactatc aaaaaaccat acgtcccaac aaggcagtca tcaaccagg 1320
 caccagggtc aggtaccgaa tgatccacca atttcaagac cagtgttaca accatcaaca 1380
 gtcaccttgg acgtgttccg tgggtactgt agacagacat ttggaaatcg atttttgaa 1440
 ggagaaaatgt aacaatccgg cgcaataatt cggtctagta acaaattcat tgaagaattt 1500
 gattcgccga tttgtgggt gacagttgtt cgaccgcgg tgacagacgg tgaggtttg 1560
 gagaacatca tgccggaaga tgcaccatcat gcaagttcat tttgaggctc 1620
 acatcagaaa gtgtacttt ctcaggagag gggccagaag tttagtattt gaacgaaaaa 1680
 tggggaaacaa ttgtgtacta tgagaaaaat ttgcaaattt gcgagaaaaa atgttcgaga 1740
 gggaaatttcc acgtggatgg cggattcatt tgctctgaga atcggttacag tctcgactt 1800
 gagccaaatc caatttagaga accagtggcg tttaaagttc gtaaagcaat agtggatgga 1860
 attcgcttt cctacaaaaa agacgggagt gtttggcttcaaaaacccat gaagtacccg 1920
 gtatttgcata cttctggta tctcgacgag caatcaggag gcctaaagaaa ggataaagt 1980
 cacaaagttt acggatgtgc gtctatcaaa acgtttggct tcaacgttcaaaaccaatc 2040
 atcagagacg cgcttcttc caagcaatg gcaacaatgt acttgcaagg aaaattgact 2100
 ccgatgaatt atatctacga gaagaagact caggaagagc tgcgaaggga agcaacacgc 2160
 accactgatt cattggccaa gtactgttgc gtccgtgtct cgttctgcaaa aggatttgg 2220
 gaagcatacc cagaacgcggc gtcaattcat gattgtccag tttggatttga gttgaaaatc 2280
 aacattgcct acgatttcat ggattcaatc tgccagtaca taaccaactg cttcgagccg 2340
 cttaggaatgg aagattttgc aaaattggaa atcaacgtca gtgatgacta aatgataact 2400
 ttttcactc accctactag atactgattt agtcttattt caaatcatcc aacgatatca 2460
 aacttttcc tttgaactttt gcatactatg ttatcacaag ttccaagcag tttcaataca 2520
 aacataggat atgttaacaa ctttgataaa gaatcaagttt accaactgtt cattgtgagc 2580
 tttgagctgt atagaaggac aatgtatccc atacctaattt cttaatagt catcagtcac 2640
 tggccccca ccaattttt cgattcgcat atgtcatata ttgcaccgtg gcccctttta 2700
 ttgttaactttt taatataattt tcttcccaac ttgtgaatattt gattgtatgaa ccaccatttt 2760
 gagtaataaaa tgtatTTT gtgg 2784

<210> 40

<211> 796

<212> PRT

<213> *Caenorhabditis elegans*

<400> 40

Met	Lys	Leu	Ile	Ala	Thr	Ser	Leu	Leu	Val	Pro	Asp	Glu	His	Thr	Pro
1															15
Met	Met	Ser	Pro	Val	Asn	Thr	Thr	Thr	Lys	Ile	Leu	Gln	Arg	Ser	Gly
															20
Ile	Lys	Met	Glu	Ile	Pro	Pro	Tyr	Leu	Asp	Pro	Asp	Ser	Gln	Asp	Asp
															35
Asp	Pro	Glu	Asp	Gly	Val	Asn	Tyr	Pro	Asp	Pro	Asp	Leu	Phe	Asp	Thr
															50
Lys	Asn	Thr	Asn	Met	Thr	Glu	Tyr	Asp	Leu	Asp	Val	Leu	Lys	Leu	Gly
															65
Lys	Pro	Ala	Val	Asp	Glu	Ala	Arg	Lys	Lys	Ile	Glu	Val	Pro	Asp	Ala
															85
Ser	Ala	Pro	Pro	Asn	Lys	Ile	Val	Glu	Tyr	Leu	Met	Tyr	Tyr	Arg	Thr
															100
Leu	Lys	Glu	Ser	Glu	Leu	Ile	Gln	Leu	Asn	Ala	Tyr	Arg	Thr	Lys	Arg
															115
Asn	Arg	Leu	Ser	Leu	Asn	Leu	Val	Lys	Asn	Asn	Ile	Asp	Arg	Glu	Phe
															130
Asp	Gln	Lys	Ala	Cys	Glu	Ser	Leu	Val	Lys	Lys	Leu	Lys	Asp	Lys	Lys
															145
Asn	Asp	Leu	Gln	Asn	Leu	Ile	Asp	Val	Val	Leu	Ser	Lys	Gly	Thr	Lys
															165
Tyr	Thr	Gly	Cys	Ile	Thr	Ile	Pro	Arg	Thr	Leu	Asp	Gly	Arg	Leu	Gln
															170
															175

200
190
180
170
160
150
140
130
120
110
100
90
80
70
60
50
40
30
20
10
P

180	185	190
Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu Trp		
195	200	205
Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His Cys		
210	215	220
Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro Tyr		
225	230	235
240		
His Tyr Glu Ile Val Ile Gly Thr Met Ile Val Gly Gln Arg Asp His		
245	250	255
Asp Asn Arg Asp Met Pro Pro His Gln Arg Tyr His Thr Pro Gly		
260	265	270
Arg Gln Asp Pro Val Asp Asp Met Ser Arg Phe Ile Pro Pro Ala Ser		
275	280	285
Ile Arg Pro Pro Met Asn Met His Thr Arg Pro Gln Pro Met Pro		
290	295	300
Gln Gln Leu Pro Ser Val Gly Ala Thr Phe Ala His Pro Leu Pro His		
305	310	315
320		
Gln Ala Pro His Asn Pro Gly Val Ser His Pro Tyr Ser Ile Ala Pro		
325	330	335
Gln Thr His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln		
340	345	350
Met Pro Gln Met Pro Pro Pro Leu His Gln Gly Tyr Gly Met Asn Gly		
355	360	365
Pro Ser Cys Ser Ser Glu Asn Asn Asn Pro Phe His Gln Asn His His		
370	375	380
Tyr Asn Asp Ile Ser His Pro Asn His Tyr Ser Tyr Asp Cys Gly Pro		
385	390	395
400		
Asn Leu Tyr Gly Phe Pro Thr Pro Tyr Pro Asp Phe His His Pro Phe		
405	410	415
Asn Gln Gln Pro His Gln Pro Pro Gln Leu Ser Gln Asn His Thr Ser		
420	425	430
Gln Gln Gly Ser His Gln Pro Gly His Gln Gly Gln Val Pro Asn Asp		
435	440	445
Pro Pro Ile Ser Arg Pro Val Leu Gln Pro Ser Thr Val Thr Leu Asp		
450	455	460
Val Phe Arg Arg Tyr Cys Arg Gln Thr Phe Gly Asn Arg Phe Phe Glu		
465	470	475
480		
Gly Glu Ser Glu Gln Ser Gly Ala Ile Ile Arg Ser Ser Asn Lys Phe		
485	490	495
Ile Glu Glu Phe Asp Ser Pro Ile Cys Gly Val Thr Val Val Arg Pro		
500	505	510
515		
Arg Met Thr Asp Gly Glu Val Leu Glu Asn Ile Met Pro Glu Asp Ala		
525		
Pro Tyr His Asp Ile Cys Lys Phe Ile Leu Arg Leu Thr Ser Glu Ser		
530	535	540
540		
Val Thr Phe Ser Gly Glu Gly Pro Glu Val Ser Asp Leu Asn Glu Lys		
545	550	555
560		
Trp Gly Thr Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys		
565	570	575
580		
Lys Cys Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser		
585		590
Glu Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro		
595	600	605
610		
Val Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser		
615		620
Tyr Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro		
625	630	635
640		
Val Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys		

645	650	655
Lys Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe		
660	665	670
Gly Phe Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys		
675	680	685
Gln Met Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr		
690	695	700
Ile Tyr Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg		
705	710	715
Thr Thr Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys		
725	730	735
Lys Gly Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys		
740	745	750
Pro Val Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp		
755	760	765
Ser Ile Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu		
770	775	780
Asp Phe Ala Lys Leu Gly Ile Asn Val Ser Asp Asp		
785	790	795

<210> 41
<211> 858
<212> PRT
<213> Caenorhabditis elegans

<400> 41

Met Gly Asp His His Asn Leu Thr Gly Leu Pro Gly Thr Ser Ile Pro		
1	5	10
Pro Gln Phe Asn Tyr Ser Gln Pro Gly Thr Ser Thr Gly Gly Pro Leu		
20	25	30
Tyr Gly Gly Lys Pro Ser His Gly Leu Glu Asp Ile Pro Asp Val Glu		
35	40	45
Glu Tyr Glu Arg Asn Leu Leu Gly Ala Gly Ala Phe Asn Leu Leu		
50	55	60
Asn Val Gly Asn Met Ala Asn Val Pro Asp Glu His Thr Pro Met Met		
65	70	75
Ser Pro Val Asn Thr Thr Thr Lys Ile Leu Gln Arg Ser Gly Ile Lys		
85	90	95
Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp Asp Pro		
100	105	110
Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr Lys Asn		
115	120	125
Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly Lys Pro		
130	135	140
Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala Ser Ala		
145	150	155
Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr Leu Lys		
165	170	175
Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg Asn Arg		
180	185	190
Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe Asp Gln		
195	200	205
Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys Asn Asp		
210	215	220
Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys Tyr Thr		
225	230	235
Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln Val His		

1000
900
800
700
600
500
400
300
200
100

245	250	255
Gly Arg Lys Gly Phe Pro His Val Val Tyr	Gly Lys Leu Trp Arg Phe	
260	265	270
Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His Cys Lys His		
275	280	285
Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro Tyr His Tyr		
290	295	300
Glu Ile Val Ile Gly Thr Met Ile Val Gly Gln Arg Asp His Asp Asn		
305	310	315
Arg Asp Met Pro Pro His Gln Arg Tyr His Thr Pro Gly Arg Gln		
325	330	335
Asp Pro Val Asp Asp Met Ser Arg Phe Ile Pro Pro Ala Ser Ile Arg		
340	345	350
Pro Pro Pro Met Asn Met His Thr Arg Pro Gln Pro Met Pro Gln Gln		
355	360	365
Leu Pro Ser Val Gly Ala Thr Phe Ala His Pro Leu Pro His Gln Ala		
370	375	380
Pro His Asn Pro Gly Val Ser His Pro Tyr Ser Ile Ala Pro Gln Thr		
385	390	395
His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln Met Pro		
405	410	415
Gln Met Pro Pro Leu His Gln Gly Tyr Gly Met Asn Gly Pro Ser		
420	425	430
Cys Ser Ser Glu Asn Asn Asn Pro Phe His Gln Asn His His Tyr Asn		
435	440	445
Asp Ile Ser His Pro Asn His Tyr Ser Tyr Asp Cys Gly Pro Asn Leu		
450	455	460
Tyr Gly Phe Pro Thr Pro Tyr Pro Asp Phe His His Pro Phe Asn Gln		
465	470	475
Gln Pro His Gln Pro Pro Gln Leu Ser Gln Asn His Thr Ser Gln Gln		
485	490	495
Gly Ser His Gln Pro Gly His Gln Gly Gln Val Pro Asn Asp Pro Pro		
500	505	510
Ile Ser Arg Pro Val Leu Gln Pro Ser Thr Val Thr Leu Asp Val Phe		
515	520	525
Arg Arg Tyr Cys Arg Gln Thr Phe Gly Asn Arg Phe Phe Glu Gly Glu		
530	535	540
Ser Glu Gln Ser Gly Ala Ile Ile Arg Ser Ser Asn Lys Phe Ile Glu		
545	550	555
Glu Phe Asp Ser Pro Ile Cys Gly Val Thr Val Val Arg Pro Arg Met		
565	570	575
Thr Asp Gly Glu Val Leu Glu Asn Ile Met Pro Glu Asp Ala Pro Tyr		
580	585	590
His Asp Ile Cys Lys Phe Ile Leu Arg Leu Thr Ser Glu Ser Val Thr		
595	600	605
Phe Ser Gly Glu Gly Pro Glu Val Ser Asp Leu Asn Glu Lys Trp Gly		
610	615	620
Thr Ile Val Tyr Tyr Glu Asn Leu Gln Ile Gly Glu Lys Lys Cys		
625	630	635
Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser Glu Asn		
645	650	655
Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val Ala		
660	665	670
Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr Lys		
675	680	685
Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val Phe		
690	695	700
Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys Asp		

705	710	715	720
Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe Gly Phe			
725	730	735	
Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys Gln Met			
740	745	750	
Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr Ile Tyr			
755	760	765	
Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg Thr Thr			
770	775	780	
Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly			
785	790	795	800
Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys Pro Val			
805	810	815	
Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp Ser Ile			
820	825	830	
Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu Asp Phe			
835	840	845	
Ala Lys Leu Gly Ile Asn Val Ser Asp Asp			
850	855		

<210> 42
<211> 892
<212> PRT
<213> Caenorhabditis elegans

<400> 42			
Met Gly Asp His His Asn Leu Thr Gly Leu Pro Gly Thr Ser Ile Pro			
1	5	10	15
Pro Gln Phe Asn Tyr Ser Gln Pro Gly Thr Ser Thr Gly Gly Pro Leu			
20	25	30	
Tyr Gly Gly Lys Pro Ser His Gly Leu Glu Asp Ile Pro Asp Val Glu			
35	40	45	
Glu Tyr Glu Arg Asn Leu Leu Gly Ala Gly Ala Gly Phe Asn Leu Leu			
50	55	60	
Asn Val Gly Asn Met Ala Asn Glu Phe Lys Pro Ile Ile Thr Leu Asp			
65	70	75	80
Thr Lys Pro Pro Arg Asp Ala Asn Lys Ser Leu Ala Phe Asn Gly Gly			
85	90	95	
Leu Lys Leu Ile Thr Pro Lys Thr Glu Val Pro Asp Glu His Thr Pro			
100	105	110	
Met Met Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly			
115	120	125	
Ile Lys Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp			
130	135	140	
Asp Pro Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr			
145	150	155	160
Lys Asn Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly			
165	170	175	
Lys Pro Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala			
180	185	190	
Ser Ala Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr			
195	200	205	
Leu Lys Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg			
210	215	220	
Asn Arg Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe			
225	230	235	240
Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys			

EQUUS 2000

245	250	255
Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys		
260	265	270
Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln		
275	280	285
Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu Trp		
290	295	300
Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His Cys		
305	310	315
Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro Tyr		
325	330	335
His Tyr Glu Ile Val Ile Gly Thr Met Ile Val Gly Gln Arg Asp His		
340	345	350
Asp Asn Arg Asp Met Pro Pro His Gln Arg Tyr His Thr Pro Gly		
355	360	365
Arg Gln Asp Pro Val Asp Asp Met Ser Arg Phe Ile Pro Pro Ala Ser		
370	375	380
Ile Arg Pro Pro Pro Met Asn Met His Thr Arg Pro Gln Pro Met Pro		
385	390	395
Gln Gln Leu Pro Ser Val Gly Ala Thr Phe Ala His Pro Leu Pro His		
405	410	415
Gln Ala Pro His Asn Pro Gly Val Ser His Pro Tyr Ser Ile Ala Pro		
420	425	430
Gln Thr His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln		
435	440	445
Met Pro Gln Met Pro Pro Pro Leu His Gln Gly Tyr Gly Met Asn Gly		
450	455	460
Pro Ser Cys Ser Ser Glu Asn Asn Asn Pro Phe His Gln Asn His His		
465	470	475
Tyr Asn Asp Ile Ser His Pro Asn His Tyr Ser Tyr Asp Cys Gly Pro		
485	490	495
Asn Leu Tyr Gly Phe Pro Thr Pro Tyr Pro Asp Phe His His Pro Phe		
500	505	510
Asn Gln Gln Pro His Gln Pro Pro Gln Leu Ser Gln Asn His Thr Ser		
515	520	525
Gln Gln Gly Ser His Gln Pro Gly His Gln Gly Gln Val Pro Asn Asp		
530	535	540
Pro Pro Ile Ser Arg Pro Val Leu Gln Pro Ser Thr Val Thr Leu Asp		
545	550	555
Val Phe Arg Arg Tyr Cys Arg Gln Thr Phe Gly Asn Arg Phe Phe Glu		
565	570	575
Gly Glu Ser Glu Gln Ser Gly Ala Ile Ile Arg Ser Ser Asn Lys Phe		
580	585	590
Ile Glu Glu Phe Asp Ser Pro Ile Cys Gly Val Thr Val Val Arg Pro		
595	600	605
Arg Met Thr Asp Gly Glu Val Leu Glu Asn Ile Met Pro Glu Asp Ala		
610	615	620
Pro Tyr His Asp Ile Cys Lys Phe Ile Leu Arg Leu Thr Ser Glu Ser		
625	630	635
Val Thr Phe Ser Gly Glu Gly Pro Glu Val Ser Asp Leu Asn Glu Lys		
645	650	655
Trp Gly Thr Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys		
660	665	670
Lys Cys Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser		
675	680	685
Glu Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro		
690	695	700
Val Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser		

EQUATION

705	710	715	720
Tyr Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro			
725	730	735	
Val Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys			
740	745	750	
Lys Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe			
755	760	765	
Gly Phe Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys			
770	775	780	
Gln Met Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr			
785	790	795	800
Ile Tyr Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg			
805	810	815	
Thr Thr Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys			
820	825	830	
Lys Gly Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys			
835	840	845	
Pro Val Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp			
850	855	860	
Ser Ile Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu			
865	870	875	880
Asp Phe Ala Lys Leu Gly Ile Asn Val Ser Asp Asp			
885	890		

<210> 43

<211> 3499

<212> DNA

<213> Caenorhabditis elegans

<400> 43

tatctttca agccgaagca atcaagacct caaagccaat caactctact cactttctt 60
cagaaccta acttttgtg tcactttccc caaaaaccgt tcaagctgc gccttcactc 120
tcatccccctc ctcttactcc ttctttctcg tccgctacta ctgtatcttc tggacatcta 180
cctgtataca caccagtggc cagtcatctg ccattacaat ttcatcaatt gacacttctt 240
caacaacaac cgccgtcctc attcaactccc gattcttctt catcctcaac atcgtcgtct 300
ttggctgaaa ttccgaaaga cgttatgtg gagatgtctgg tagatcaggaa aactgtatgca 360
tcgtcatccg cttcacgtc cacccatctt gtttccatgtt tcggagcggaa cacgttcatg 420
aatacaccgg atgatgtatgat gatgaatgtat gatatggaaac cgatttcctcg tgatcggtgc 480
aatacgtggc caatcgtag gcccgaactc gaaccaccac tcaactcgag tcccattatt 540
catgaacaaa ttccgtaaaga agatgtctgc ctatacgggaa gcaatgagca atgtggacag 600
ctcggcggag catcttcaa cgggtcgac gcaatgttcc atactccaga tggaaagcaat 660
tctcatcaga catcggttcc tggagtttc agaatgttcc atcgccaga cgataccgtta 720
tcggggaaaaa agacaacgac cagacggaaac gcttggggaa atatgtcata tgctgaactt 780
atcaactacag ccattatggc tagtccagag aaacggttaa ctcttgccaca agtttacgaa 840
tggatggttcc agaatgttcc atacttcagg gataaggagat ttcgaaacag ttcagctggaa 900
tggaaagaact cgatccgtca caatctgttctt cttccatgttgc aattcagaat 960
gaaggagccg gaaagagctc gtgggtgggtt attaatccag atgcaaaaggcc aggaatgaat 1020
ccacggcgta cacgtgaacg atccaataact attgagacga ctacaaaggcc tcaactcgaa 1080
aaatctcgcc gcggagccaa gaagaggata aaggagagag cattgtatggg ctcccttcac 1140
tcgacactta atggaaattt gattgccga tcgatcaaa cgatttctca cgatttgtat 1200
gatgatgatc aatgcaagga gcatttgcata acgttccatc atctttccgt ccccgaaactc 1260
aatcgaaacct ctcgatttccgtt gatcgtcgat ctcgttcc tccagctatt ggaagtgcata 1320
tctatgtatca tctagaattt ccatcatggg ttggcgaatc ggttcccgaa attccaatgt 1380
atattgttca tagaactgtat caaatgcgtt tcgatgcac tactcatagt tggtgagtt 1440
cagattaagc aggagtcgaa gccgattaaac acggaaccaa ttgctccacc accatcatac 1500
cacgagttga acagtgtccg tggatgtgtt gtcagaatc cacttcttgc aaatccaatt 1560
gtgccaagca ctaacttcaa gccaatgcac ctaccgggtt cctatggaaa ctatcaaaat 1620

gggtggaaataa	ctccaatcaa	ttggctatca	acatccaaact	catctccact	gcctggaatt	1680
caatcggttg	gaatttgtac	tgcacagcat	actgtcgctt	cttcatcgcc	tcttcaatt	1740
gatttggaaa	atctgacact	tccccgatcg	ccactgatgg	atactatgga	tgttgatgca	1800
ttgatcagac	atgagcttag	tcaagctgga	gggcagcata	ttcattttga	tttgtaaatt	1860
ctcttcattt	tgttcccct	ggtgttggc	gaaagagaga	tagcaaagca	gcgaggagtg	1920
aggttaagcag	caataaaaaat	tttggatttt	tttttgggtt	ttccagaaaat	aatcgatttt	1980
ctggaaaatt	tcaaaaaaaaa	atcggaaattt	ttagttaaatt	atttgatgag	aaaaaaaaat	2040
tagaaaacat	aaggaaaaat	gaaaagcgtt	tttttttttc	gaaaatttta	gaattctcct	2100
acatttccaa	taagggcctt	agaactgcaa	acaaaacaaaa	attggaattt	tcgaatcaa	2160
aagtcccg	ataaaaagtag	ttcgaatatt	aaaaagcatt	taatttcctc	ttaaaaaaaaa	2220
ttgaataata	gccgaaaattt	gcagattttt	ttctgaaaa	tcgaaaaacc	aaaatttttt	2280
gattttttaa	attttttttt	tactttccag	atagaaaaat	cattagcact	gaaaattatt	2340
tgaaaaaaaaa	cttcaaatac	aaattttgtt	ttcgaaaaaaaa	aaaattttaa	tatataatttt	2400
cagaaaatctt	ccgtcttcat	cttttcaaat	ccctacctac	acacactcaa	cgatcatcac	2460
agccagacca	tcaatattct	tccaaatttt	gacgtcgta	attttttttc	agtttttca	2520
aaaactctat	tttctatttt	ctgtcgttt	ttccccccttc	tctcgctcaa	ttccaacaca	2580
ttcatcccg	tgacgtcg	taataataat	ataaaatacc	tcttctctt	ttcttccct	2640
aatgcgaat	atcgaaaaac	cgttgattat	tacctctttt	ttcttggttt	ttttttctct	2700
ctctctctcc	cgtcatccag	gttcttact	ctttaaatgc	tacctctatc	ccatcttttt	2760
cgctgtaaat	ttgtttcgca	atcaaaaactg	ctaaaacaca	ttccccaatc	tgttttttt	2820
aattgaattt	ttcaaaaaat	ttgatttctt	gatttcttctt	gtaattctt	aattttcctc	2880
tttttttcc	ccctggtagc	aaatgtctag	cgatttctt	tctttttttt	ttaactttc	2940
acatctggcc	gattcgaatc	ctccgtatac	acacacacat	agtaatctac	ctccaaaatt	3000
ttactgaaag	atgtgatccc	ctctctgtct	ccctctacaa	aacattattt	gtctgtttgt	3060
gtatattgcc	accacgtcg	ttttaaatta	aaaccatcgt	tttttcttct	tttctacttt	3120
tttctcgaaa	aatttaacaa	cacacaaaaa	aatcctcaa	aaaatctcag	ttttaaatgg	3180
tgtggcaata	tatcgatcc	ccctctacac	cagaacagtc	ttgcaatttc	agagaatgat	3240
tttcagattt	ttcatatcac	aggccccctt	ttttgcttg	ttttttctc	tacctcttct	3300
tcttttcat	ctatttctct	ctcttggttt	ctctctgtta	tcctgtacat	tttccttcca	3360
atttttctg	gctatttctg	attttcgagt	tcatattctc	taacgtctcac	tttctctcg	3420
gccacgcccc	cttttctgc	tccctccgcc	cccaaataata	tttgcgactg	tatgatgatg	3480
atqatqattt	aataaaaaat					3499

<210> 44
<211> 2704
<212> DNA
<213> *Caenorhabditis elegans*

```

<400> 44
ttacacgtgg ccaatgcaac aatacatcta tcaggaatcg tcagcaacca ttccccatca 60
ccatttaaat caacacaaca atccgtatca tccaatgcat cctcatcatc aattacctca 120
tatgcaacaa ctccctcaac ctctattgaa tcttaacatg acgacgttaa catcttctgg 180
cagttccgtg gccagttcca ttggaggcgg agctcaatgc tctccgtcc cgtcgggctc 240
ctcgaccgct gcaacaaaatt cctctcaaca gcagcagacc gttggtaaa tgcttgctgc 300
atcggtgcct tgttcttcat ctggcatgac acttggaaatg tcacttaatc tgtcacaagg 360
cggtggtcca atgcccggcaa aaaagaagcg ttgtcgtaag aagccaaaccc atcaattggc 420
acagaagaaa ccgaatccat ggggtgagga atccatttcg gatatcatcg ccaaagcatt 480
ggaatcgcg ccagacggaa ggcttaaact caatgagatt tatcaatggt tctctgataa 540
tattccctac tttggagaac gatcttagtcc cgaggaggcc gccggatgga agaactcgat 600
ccgtcacaat ctgtcttcc attctcggtt catgcaatt cagaatgaag gagccggaaa 660
gagctcggtg tgggttatta atccagatgc aaagccagga atgaatccac ggcgtacacg 720
tgaacgatcc aatactattg agacgactac aaaggctcaa ctcgaaaaat ctcggccgg 780
agccaagaag aggataaagg agagagcatt gatgggctcc cttcaactcgaa cacttaatgg 840
aaattcgatt gcccggatcg ttccaaacgat ttctcacgat ttgtatgatg atgattcaat 900
gcaaggagca tttgtataacg ttccatcatc tttccgtccc cgaactcaat cgaacctctc 960
gattccctgg tcgtcgatctc gtgtttctcc agctattgga agtgtatatct atgtatgatct 1020
agaattccca tcatgggttg gcgaaatcggt tccagcaatt ccaagtgata ttgttgatag 1080
aactgatcaa atgcgtatcg atgcaactac tcatattggg ggagttcaga ttaagcagga 1140

```

gtcgaagccg attaagacgg aaccaattgc tccaccacca tcataccacg agttgaacag 1200
 tgtccgtgaa tcgtgtgctc agaatccact tcttcgaaat ccaattgtgc caagcactaa 1260
 cttcaagcca atgccactac cgggtgccta tggaaactat caaaatggtg gaataactcc 1320
 aatcaattgg ctatcaacat ccaactcatc tccactgcct ggaattcaat cgtgtggaat 1380
 tgttagctgca cagcatactg tcgcttctc atcgctctt ccaattgatt tggaaaatct 1440
 gacacttccc gatcagccac tgatggatac tatggatgtt gatgcattga tcagacatga 1500
 gctgagtcaa gctggaggc acgatattca tttgatttg taaaattctct tcattttgtt 1560
 tccccctgggt ttgttcgaaa gagagatagc aaagcagcga ggagtggaaa atctccgtc 1620
 ttcatcttt caaatcccta cctacacaca ctcaacgatc atcacagcca gaccatcaat 1680
 attcttccaa attttgacgt cgtaatttt tttttagt tttcaaaaac tctattttct 1740
 attttctgtc gtttgttccc ctttctctcg tctaattcca acacattcat cccagtgacg 1800
 tcgtgtaata ataataaaaa atacctctc tctcttctt cccctaattgc gaaatatcga 1860
 aaaaccgttg attattacct cttttttctt gtttttttt tctctctctc tctccgtca 1920
 tccaggttct tcactctta aatgctacct ctatcccattt ttttcgctg taaaattgtt 1980
 tcgcaatcaa aactgctaaa acacattccc caatctgtct ttttaattt aattttcaa 2040
 aaaatttgat ttcttgattt ctcttgtaat tcttaattt tcctctttt ttcccccctg 2100
 gtagcaaatg tctagcgatt ctcttctt tttgtttaa cttcacatc tggccgattc 2160
 gaatccctccg tatacacaca cacatagtaa tctacctcca aaattttact gaaagatgtg 2220
 atcccccttc tgcctccctc tacaaaacat tatttgcctg ttgtgtata ttgcaccac 2280
 gtcgatttta aattaaaaacc atcgaaaaat cttcttttctt actttttctt cgaaaaattt 2340
 aacaacacac aaaaaaaaaatcc tcaaaaaat ctcagttta aatgggtgtgg caatataatcg 2400
 gatccccctc tacaccagaa cagtcgtca atttcagaga atgattttca gattttcat 2460
 atcacaggcc ccctttttt gttgtttt ttctctacct ctctttctt tcatttttatt 2520
 tctctctctt gttttctctc tggttatccctg tacattttcc ttccaattct ttctggctat 2580
 ttctgatttt cgagttcata ttctctacgt ctcaatttctt ctgcgcac gccccctttt 2640
 tcgtctccct ccgcccccaa atatattgc gactgtatga tgatgtatgat gatttaataa 2700
 aaat 2704

<210> 45

<211> 510

<212> PRT

<213> Caenorhabditis elegans

<400> 45

Met	Met	Glu	Met	Lle	Val	Asp	Gln	Gly	Thr	Asp	Ala	Ser	Ser	Ser	Ala
1									10						15
Ser	Thr	Ser	Thr	Ser	Ser	Val	Ser	Arg	Phe	Gly	Ala	Asp	Thr	Phe	Met
									25						30
Asn	Thr	Pro	Asp	Asp	Val	Met	Met	Asn	Asp	Asp	Met	Glu	Pro	Ile	Pro
									35	40					45
Arg	Asp	Arg	Cys	Asn	Thr	Trp	Pro	Met	Arg	Arg	Pro	Gln	Leu	Glu	Pro
									50	55					60
Pro	Leu	Asn	Ser	Ser	Pro	Ile	Ile	His	Glu	Gln	Ile	Pro	Glu	Glu	Asp
									65	70					80
Ala	Asp	Leu	Tyr	Gly	Ser	Asn	Glu	Gln	Cys	Gly	Gln	Leu	Gly	Gly	Ala
									85	90					95
Ser	Ser	Asn	Gly	Ser	Thr	Ala	Met	Leu	His	Thr	Pro	Asp	Gly	Ser	Asn
									100	105					110
Ser	His	Gln	Thr	Ser	Phe	Pro	Ser	Asp	Phe	Arg	Met	Ser	Glu	Ser	Pro
									115	120					125
Asp	Asp	Thr	Val	Ser	Gly	Lys	Lys	Thr	Thr	Arg	Arg	Asn	Ala	Trp	
									130	135					140
Gly	Asn	Met	Ser	Tyr	Ala	Glu	Leu	Ile	Thr	Thr	Ala	Ile	Met	Ala	Ser
									145	150					160
Pro	Glu	Lys	Arg	Leu	Thr	Leu	Ala	Gln	Val	Tyr	Glu	Trp	Met	Val	Gln
									165	170					175
Asn	Val	Pro	Tyr	Phe	Arg	Asp	Lys	Gly	Asp	Ser	Asn	Ser	Ser	Ala	Gly
									180	185					190

© 2007-2008

Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met
195 200 205
Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn
210 215 220
Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser
225 230 235 240
Asn Thr Ile Glu Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg
245 250 255
Gly Ala Lys Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His
260 265 270
Ser Thr Leu Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser
275 280 285
His Asp Leu Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val
290 295 300
Pro Ser Ser Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly
305 310 315 320
Ser Ser Ser Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp
325 330 335
Leu Glu Phe Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser
340 345 350
Asp Ile Val Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His
355 360 365
Ile Gly Gly Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu
370 375 380
Pro Ile Ala Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly
385 390 395 400
Ser Cys Ala Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr
405 410 415
Asn Phe Lys Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn
420 425 430
Gly Gly Ile Thr Pro Ile Asn Trp Leu Ser Thr Ser Asn Ser Ser Pro
435 440 445
Leu Pro Gly Ile Gln Ser Cys Gly Ile Val Ala Ala Gln His Thr Val
450 455 460
Ala Ser Ser Ser Ala Leu Pro Ile Asp Leu Glu Asn Leu Thr Leu Pro
465 470 475 480
Asp Gln Pro Leu Met Asp Thr Met Asp Val Asp Ala Leu Ile Arg His
485 490 495
Glu Leu Ser Gln Ala Gly Gly Gln His Ile His Phe Asp Leu
500 505 510

<210> 46
<211> 509
<212> PRT
<213> Caenorhabditis elegans

<400> 46
Met Gln Gln Tyr Ile Tyr Gln Glu Ser Ser Ala Thr Ile Pro His His
1 5 10 15
His Leu Asn Gln His Asn Asn Pro Tyr His Pro Met His Pro His His
20 25 30
Gln Leu Pro His Met Gln Gln Leu Pro Gln Pro Leu Leu Asn Leu Asn
35 40 45
Met Thr Thr Leu Thr Ser Ser Gly Ser Ser Val Ala Ser Ser Ile Gly
50 55 60
Gly Gly Ala Gln Cys Ser Pro Cys Ala Ser Gly Ser Ser Thr Ala Ala
65 70 75 80

DRAFT 6.0

Thr Asn Ser Ser Gln Gln Gln Thr Val Gly Gln Met Leu Ala Ala
85 90 95
Ser Val Pro Cys Ser Ser Ser Gly Met Thr Leu Gly Met Ser Leu Asn
100 105 110
Leu Ser Gln Gly Gly Gly Pro Met Pro Ala Lys Lys Lys Arg Cys Arg
115 120 125
Lys Lys Pro Thr Asp Gln Leu Ala Gln Lys Lys Pro Asn Pro Trp Gly
130 135 140
Glu Glu Ser Tyr Ser Asp Ile Ile Ala Lys Ala Leu Glu Ser Ala Pro
145 150 155 160
Asp Gly Arg Leu Lys Leu Asn Glu Ile Tyr Gln Trp Phe Ser Asp Asn
165 170 175
Ile Pro Tyr Phe Gly Glu Arg Ser Ser Pro Glu Glu Ala Ala Gly Trp
180 185 190
Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met Arg
195 200 205
Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro
210 215 220
Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser Asn
225 230 235 240
Thr Ile Glu Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg Gly
245 250 255
Ala Lys Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His Ser
260 265 270
Thr Leu Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser His
275 280 285
Asp Leu Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val Pro
290 295 300
Ser Ser Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly Ser
305 310 315 320
Ser Ser Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp Leu
325 330 335
Glu Phe Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser Asp
340 345 350
Ile Val Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His Ile
355 360 365
Gly Gly Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu Pro
370 375 380
Ile Ala Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly Ser
385 390 395 400
Cys Ala Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr Asn
405 410 415
Phe Lys Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn Gly
420 425 430
Gly Ile Thr Pro Ile Asn Trp Leu Ser Thr Ser Asn Ser Ser Pro Leu
435 440 445
Pro Gly Ile Gln Ser Cys Gly Ile Val Ala Ala Gln His Thr Val Ala
450 455 460
Ser Ser Ser Ala Leu Pro Ile Asp Leu Glu Asn Leu Thr Leu Pro Asp
465 470 475 480
Gln Pro Leu Met Asp Thr Met Asp Val Asp Ala Leu Ile Arg His Glu
485 490 495
Leu Ser Gln Ala Gly Gly Gln His Ile His Phe Asp Leu
500 505

<210> 47
<211> 3504

<212> DNA

<213> *Caenorhabditis elegans*

<400> 47

ttcaaaaacgt tatgcgtcga agcctacgaa gtaatgtgga ataatcgaga tttgttcgtt 3300
tccttggcca ccttgatgct cggaatggag ttgcctgagc tgtcgacgaa agcggattt 3360
gatcatttga agaaaaccct cttctgcaat ggagaaaagca aagaagaagc gagaaagttt 3420
ttcgctggaa tctacgaaga agccttcaat ggatcatggt ctacaaaaac gaattggctc 3480
ttccacgcag tcaaacacta ctga 3504

<210> 48
<211> 1167
<212> PRT
<213> Caenorhabditis elegans

<400> 48
Arg Lys Pro Trp Ser Ser Arg Ser Asp Cys Trp Thr Arg Thr Glu Leu
1 5 10 15
Arg Arg Ile Ser Gln Met His Val Asn Ile Leu His Pro Gln Leu Gln
20 25 30
Thr Met Val Glu Gln Trp Gln Met Arg Glu Arg Pro Ser Leu Glu Thr
35 40 45
Glu Asn Gly Lys Gly Ser Leu Leu Leu Asn Glu Gly Val Ala Asp
50 55 60
Ile Ile Thr Met Cys Pro Phe Gly Glu Val Ile Ser Val Val Phe Pro
65 70 75 80
Trp Phe Leu Ala Asn Val Arg Thr Ser Leu Glu Ile Lys Leu Ser Asp
85 90 95
Phe Lys His Gln Leu Phe Glu Leu Ile Ala Pro Met Lys Trp Gly Thr
100 105 110
Tyr Ser Val Lys Pro Gln Asp Tyr Val Phe Arg Gln Leu Asn Asn Phe
115 120 125
Gly Glu Ile Glu Val Ile Phe Asn Asp Gln Pro Leu Ser Lys Leu
130 135 140
Glu Leu His Gly Thr Phe Pro Met Leu Phe Leu Tyr Gln Pro Asp Gly
145 150 155 160
Ile Asn Arg Asp Lys Glu Leu Met Ser Asp Ile Ser His Cys Leu Gly
165 170 175
Tyr Ser Leu Asp Lys Leu Glu Ser Leu Asp Glu Glu Leu Arg Gln
180 185 190
Phe Arg Ala Ser Leu Trp Ala Arg Thr Lys Lys Thr Cys Leu Thr Arg
195 200 205
Gly Leu Glu Gly Thr Ser His Tyr Ala Phe Pro Glu Glu Gln Tyr Leu
210 215 220
Cys Val Gly Glu Ser Cys Pro Lys Asp Leu Glu Ser Lys Val Lys Ala
225 230 235 240
Ala Lys Leu Ser Tyr Gln Met Phe Trp Arg Lys Arg Lys Ala Glu Ile
245 250 255
Asn Gly Val Cys Glu Lys Met Met Lys Ile Gln Ile Glu Phe Asn Pro
260 265 270
Asn Glu Thr Pro Lys Ser Leu Leu His Thr Phe Leu Tyr Glu Met Arg
275 280 285
Lys Leu Asp Val Tyr Asp Thr Asp Asp Pro Ala Asp Glu Gly Trp Phe
290 295 300
Leu Gln Leu Ala Gly Arg Thr Thr Phe Val Thr Asn Pro Asp Val Lys
305 310 315 320
Leu Thr Ser Tyr Asp Gly Val Arg Ser Glu Leu Glu Ser Tyr Arg Cys
325 330 335
Pro Gly Phe Val Val Arg Arg Gln Ser Leu Val Leu Lys Asp Tyr Cys
340 345 350
Arg Pro Lys Pro Leu Tyr Glu Pro His Tyr Val Arg Ala His Glu Arg
355 360 365

2000 1999 1998 1997 1996

Lys Leu Ala Leu Asp Val Leu Ser Val Ser Ile Asp Ser Thr Pro Lys
370 375 380
Gln Ser Lys Asn Ser Asp Met Val Met Thr Asp Phe Arg Pro Thr Ala
385 390 395 400
Ser Leu Lys Gln Val Ser Leu Trp Asp Leu Asp Ala Asn Leu Met Ile
405 410 415
Arg Pro Val Asn Ile Ser Gly Phe Asp Phe Pro Ala Asp Val Asp Met
420 425 430
Tyr Val Arg Ile Glu Phe Ser Val Tyr Val Gly Thr Leu Thr Leu Ala
435 440 445
Ser Lys Ser Thr Thr Lys Val Asn Ala Gln Phe Ala Lys Trp Asn Lys
450 455 460
Glu Met Tyr Thr Phe Asp Leu Tyr Met Lys Asp Met Pro Pro Ser Ala
465 470 475 480
Val Leu Ser Ile Arg Val Leu Tyr Gly Lys Val Lys Leu Lys Ser Glu
485 490 495
Glu Phe Glu Val Gly Trp Val Asn Met Ser Leu Thr Asp Trp Arg Asp
500 505 510
Glu Leu Arg Gln Gly Gln Phe Leu Phe His Leu Trp Ala Pro Glu Pro
515 520 525
Thr Ala Asn Arg Ser Arg Ile Gly Glu Asn Gly Ala Arg Ile Gly Thr
530 535 540
Asn Ala Ala Val Thr Ile Glu Ile Ser Ser Tyr Gly Gly Arg Val Arg
545 550 555 560
Met Pro Ser Gln Gly Gln Tyr Thr Tyr Leu Val Lys His Arg Ser Thr
565 570 575
Trp Thr Glu Thr Leu Asn Ile Met Gly Asp Asp Tyr Glu Ser Cys Ile
580 585 590
Arg Asp Pro Gly Tyr Lys Lys Leu Gln Met Leu Val Lys Lys His Glu
595 600 605
Ser Gly Ile Val Leu Glu Glu Asp Glu Gln Arg His Val Trp Met Trp
610 615 620
Arg Arg Tyr Ile Gln Lys Gln Glu Pro Asp Leu Leu Ile Val Leu Ser
625 630 635 640
Glu Leu Ala Phe Val Trp Thr Asp Arg Glu Asn Phe Ser Glu Leu Tyr
645 650 655
Val Met Leu Glu Lys Trp Lys Pro Pro Ser Val Ala Ala Leu Thr
660 665 670
Leu Leu Gly Lys Arg Cys Thr Asp Arg Val Ile Arg Lys Phe Ala Val
675 680 685
Glu Lys Leu Asn Glu Gln Leu Ser Pro Val Thr Phe His Leu Phe Ile
690 695 700
Leu Pro Leu Ile Gln Ala Leu Lys Tyr Glu Pro Arg Ala Gln Ser Glu
705 710 715 720
Val Gly Met Met Leu Leu Thr Arg Ala Leu Cys Asp Tyr Arg Ile Gly
725 730 735
His Arg Leu Phe Trp Leu Leu Arg Ala Glu Ile Ala Arg Leu Arg Asp
740 745 750
Cys Asp Leu Lys Ser Glu Glu Tyr Arg Arg Ile Ser Leu Leu Met Glu
755 760 765
Ala Tyr Leu Arg Gly Asn Glu Glu His Ile Lys Ile Ile Thr Arg Gln
770 775 780
Val Asp Met Val Asp Glu Leu Thr Arg Ile Ser Thr Leu Val Lys Gly
785 790 795 800
Met Pro Lys Asp Val Ala Thr Met Lys Leu Arg Asp Glu Leu Arg Ser
805 810 815
Ile Ser His Lys Met Glu Asn Met Asp Ser Pro Leu Asp Pro Val Tyr
820 825 830

0
0
0
0
0
0
0
0
0
0
0
0
0
0

Lys Leu Gly Glu Met Ile Ile Asp Lys Ala Ile Val Leu Gly Ser Ala
835 840 845
Lys Arg Pro Leu Met Leu His Trp Lys Asn Lys Asn Pro Lys Ser Asp
850 855 860
Leu His Leu Pro Phe Cys Ala Met Ile Phe Lys Asn Gly Asp Asp Leu
865 870 875 880
Arg Gln Asp Met Leu Val Leu Gln Val Leu Glu Val Met Asp Asn Ile
885 890 895
Trp Lys Ala Ala Asn Ile Asp Cys Cys Leu Asn Pro Tyr Ala Val Leu
900 905 910
Pro Met Gly Glu Met Ile Gly Ile Ile Glu Val Val Pro Asn Cys Lys
915 920 925
Thr Ile Phe Glu Ile Gln Val Gly Thr Gly Phe Met Asn Thr Ala Val
930 935 940
Arg Ser Ile Asp Pro Ser Phe Met Asn Lys Trp Ile Arg Lys Gln Cys
945 950 955 960
Gly Ile Glu Asp Glu Lys Lys Ser Lys Lys Asp Ser Thr Lys Asn
965 970 975
Pro Ile Glu Lys Ile Asp Asn Thr Gln Ala Met Lys Lys Tyr Phe
980 985 990
Glu Ser Val Asp Arg Phe Leu Tyr Ser Cys Val Gly Tyr Ser Val Ala
995 1000 1005
Thr Tyr Ile Met Gly Ile Lys Asp Arg His Ser Asp Asn Leu Met Leu
1010 1015 1020
Thr Glu Asp Gly Lys Tyr Val His Ile Asp Phe Gly His Ile Leu Gly
1025 1030 1035 1040
His Gly Lys Thr Lys Leu Gly Ile Gln Arg Asp Arg Gln Pro Phe Ile
1045 1050 1055
Leu Thr Glu His Phe Met Thr Val Ile Arg Ser Gly Lys Ser Val Asp
1060 1065 1070
Gly Asn Ser His Glu Leu Gln Lys Phe Lys Thr Leu Cys Val Glu Ala
1075 1080 1085
Tyr Glu Val Met Trp Asn Asn Arg Asp Leu Phe Val Ser Leu Phe Thr
1090 1095 1100
Leu Met Leu Gly Met Glu Leu Pro Glu Leu Ser Thr Lys Ala Asp Leu
1105 1110 1115 1120
Asp His Leu Lys Lys Thr Leu Phe Cys Asn Gly Glu Ser Lys Glu Glu
1125 1130 1135
Ala Arg Lys Phe Phe Ala Gly Ile Tyr Glu Glu Ala Phe Asn Gly Ser
1140 1145 1150
Trp Ser Thr Lys Thr Asn Trp Leu Phe His Ala Val Lys His Tyr
1155 1160 1165

<210> 49
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Probe/primer derived from C. elegans

<400> 49
ggaaatattt taggccagat gcg

<210> 50
<211> 20
<212> DNA

23

卷之三

<213> Artificial Sequence

<220>

<223> Probe/primer derived from C. elegans

<400> 50
cgcacagtc tgaatacacc 20

<210> 51
<211> 28
<212> DNA

<213> Artificial Sequence

<220>

<223> Probe/primer derived from C. elegans

<400> 51
tctcggtt tgccgtcgga tgtctgcc 28

<210> 52
<211> 3017
<212> DNA

<213> Caenorhabditis elegans

<400> 52
gtaatcaa at tgtaaaggaa aaatattaat agtcagagta cacataaatg ggtgatcatc 60
ataatttaac gggccttcccc ggtacctcca tcccgcacaca gttcaactat tctcagcccg 120
gtaccagcac cggaggccccg ctttatggtg gaaaaccttc tcatggattt gaagatattc 180
ctgtatgtaga ggaatatgag aggaacctgc tcggggctgg agcagggttt aatctgtca 240
atgttagaaaa tatggcta at gttcccgacg agcacacacc gatgtatgtca ccagtgaata 300
caactacaaa gattctacaa cggagtggtt taaaatggta aatcccgcca tatttggatc 360
cagacagtc ggtatgtac ccggaaatgt gtgtcaacta cccggatcca gatttatttgc 420
acacaaaaaa cacaatatg accgagtacg atttggatgtt ggaaagctt gggaaaccag 480
cagtagatga agcaccggaaa aagatcgaaag ttcccgacgc tagtgcgcgc ccaaaca 540
ttttagaata ttttatgtat tatagaacgt taaaagaaag tgaactata caactgaatg 600
cgtatcgac aaaacgaaat cgattatgtc tgaacttggt caaaaacaat attgtatcgag 660
agttcgacca aaaacgttgc ggtccctgg tgaaaaattt gaaggataga aagaatgatc 720
tccagaacact gattgtatgtt gttcttcaa aaggtaaaaa atataccggt tgcattacaa 780
ttccaaggac acttgcgtgc cggttacagg tccacggaa aaaaagggttc cctcagctg 840
tctatggcaa actgtggagg ttaatggaa tgacaaaaaa cgaaacgcgt catgtggacc 900
actgcacca cgcatttgcgaa atgaaaaatgt acatgtatgt cgtgaatccc tatcaatc 960
aaatttgtcat tggaaactatg attgtgggc agagggatca tgacaatcg gatatgcgc 1020
cgcccacatca acgctaccac actccaggc ggcaggatcc agttgacgt atgagtagat 1080
ttataccacc agtccatt cgtccgcctc cgatgaacat gcacacaagg cctcagccta 1140
tgcctcaaca attgccttca gttggcgcgaa cgtttggccca tcctctccca catcaggcgc 1200
cacataaccc aggggttca catccgtact ccattgtcc acagacccat taccgggtga 1260
acatgaaccc aattccgcaa atgcccacaa tgccacaaaat gccaccacct ctccatcagg 1320
gatatggaaat gaatggccg agttgtctt cagaaaaacaa caatccattc caccaaaaatc 1380
accattataa tgatatttagc catccaaatc actattccta cgactgtggt ccgaacttgt 1440
acgggttcc aactccttat ccggattttc accatccctt caatcagca ccacaccgc 1500
cgcccacatca atcacaaaac catacgccc aacaaggcag tcatcaacca gggcaccac 1560
gtcagggtacc gaatgtatca ccaatttcaa gaccagtgtt acaaccatca acagtcaccc 1620
tggacgtgtt ccgtccgtac ttttagacaga cattttggaaa tcgatttttt gaaggagaaaa 1680
gtgaacaatc cggcgcataa attcgggtca gtaacaaattt cattgaagaa tttgtatcg 1740
cgatattgtgg tttgtacgtt gttcgaccgc ggtatgtac cgggtggatgtt ttggagaaaca 1800
tcatgcggaa agatgcacca tatcatgaca tttgtacgtt cattttggagg ctcacatcag 1860
aaagtgttaac tttctcagga gagggggccag aagttgtac tttgtacgtt aaatggggaa 1920
caattgtgtta ctatgtacgtt aatttgcaaa ttggcgagaa aaaaatgttcg agaggaaatt 1980

tccacgtgga	tggcggattc	atttgctcg	agaatcgta	cagtctcgga	cttgagccaa	2040
atccaattag	agaaccagtg	gcgtttaaag	ttcgtaaagc	aatagtggat	ggaattcgct	2100
tttcctacaa	aaaagacggg	agtgttggc	ttcaaaaccc	catgaagtagc	ccggattttg	2160
tcacttctgg	gtatctcgac	gagcaatcg	gaggcctaaa	gaaggataaa	gtgcacaaaag	2220
tttacggatg	tgcgtctatc	aaaacgttg	gcttcaacgt	ttccaaacaa	atcatcagag	2280
acgcgctct	ttccaagcaa	atggcaacaa	tgtacttgca	aggaaaattg	actccgatga	2340
attatatcta	cgagaagaag	actcaggaag	agctcgaaag	ggaagcaaca	cgcacccactg	2400
attcatttggc	caagtactgt	tgtgtccgt	tctcgttctg	caaaggattt	ggagaagcat	2460
acccagaacg	cccgtaatt	catgattgtc	cagtttgat	tgagttgaaa	atcaacattg	2520
cctacgattt	catggattca	atctgcccagt	acataaccaa	ctgcttcgag	ccgcttaggaa	2580
tggaagattt	tgcaaaaattg	ggaatcaacg	tcagtgtatga	ctaaatgata	actttttca	2640
ctcacccotac	tagatactga	tttagtctta	ttccaaatca	tccaacgata	tcaaactttt	2700
tcctttgaac	tttgatact	atgttatcac	aagttccaag	cagttcaat	acaaacatag	2760
gatatgttaa	caacttttga	taagaatcaa	gttaccaact	gttcattgtg	agctttgagc	2820
tgtatagaag	gacaatgtat	cccatatcc	aatcttaat	agtcatcagt	cactggtccc	2880
gcaccaattt	tttcgattcg	catatgtcat	atattgcacc	gtggccctt	ttattgtaac	2940
tttaaatata	tttcttccc	aacctgtgaa	tatgattgtat	gaaccacccat	tttgagtaat	3000
aaatgtatTTT	ttttgggg					3017

<210> 53

<211> 3119

<212> DNA

<213> *Caenorhabditis elegans*

<400> 53

gttaatcaaat	tgtaaaaggaa	aaataattaat	agtcaagagta	cacataaaatg	ggtgatcatc	60
ataatttaac	gggccttccc	ggtacacctca	tcccgccaca	gttcaactat	tctcagcccc	120
gtaccagcac	cggaggcccg	ctttatggtg	aaaaaaccttc	tcatggattt	gaagatattc	180
ctgatgtaga	ggaatatatgag	aggaacctgc	tcggggctgg	agcaggtttt	aatctgctca	240
atgttagaaaa	tatggctaat	gaatttaaac	caataatcac	attggacacg	aaaccacactc	300
gtgatgcca	caagtcat	gcattcaatg	gcggggtgaa	gctaattact	ccgaaaactg	360
aagttcccg	cgagcacaca	ccgatgatgt	caccagtgaa	tacaactaca	aagattctac	420
aacggagtgg	tataaaatg	gaaatccgc	catatggta	tccagacagt	caggatgatg	480
acccggaaaga	tgggtcaac	tacccggatc	cagatttatt	tgacacaaaaa	aacacaaaata	540
tgaccgaga	cgatttggat	gtgttgaagc	ttggaaaacc	agcagtagat	gaagcacgga	600
aaaagatcga	agttccgcac	gctagtgcgc	cgccaaacaa	aattgtagaa	tattttagatgt	660
attatagaac	gttaaaagaa	agtgaactca	tacaactgaa	tgcgtatcg	acaaaacgaa	720
atcgattatc	gttgaacttg	gtcaaaaaca	atattgatcg	agagttcgac	caaaaagctt	780
gchgagtcct	gggtaaaaaa	ttgaaggata	agaagaatga	tctccagaaac	ctgattgatg	840
tggttcttc	aaaaggtaca	aaatataccg	gttgcattac	aattccaagg	acacttgatg	900
gcccgttaca	ggtccacgga	agaaaaggtt	tccctcacgt	agtctatggc	aaactgtgga	960
ggtttaatga	aatgacaaaa	aacgaaacgc	gtcatgtgga	ccactgcaag	cacgcatttgc	1020
aaatgaaaag	tgacatggta	tgcgtgaatc	cctatcacta	cgaaaattgtc	attggaaacta	1080
tgatttgtgg	gcagagggat	catgacaat	gagatatgcc	gccgcacat	caacgctacc	1140
acactccagg	tgcgcggat	ccagttgacg	atatgatgt	atttatacca	ccagttcca	1200
ttcgtccgccc	tccgatgaac	atgcacacaa	ggccctcagcc	tatgcctcaa	caattgcctt	1260
cagttggcgc	aacgtttgcc	catcctctcc	cacatcaggc	gccacataac	ccaggggttt	1320
cacatcgta	ctccattgtct	ccacagaccc	attacccgtt	gaacatgaac	ccaattccgc	1380
aaatgcccga	aatgcccacaa	atgcacacac	cttccatca	gggatatgga	atgaatgggc	1440
cgagttgtc	ttcagaaaaac	aacaatccat	tccacaaaaa	tcaccattat	aatgatatta	1500
gccatccaaa	tcactattcc	tacgactgtg	gtccgaactt	gtacgggtt	ccaaactcctt	1560
atccggattt	tcaccatct	ttcaatcagc	aaccacacca	gccgccacaa	ctatcaca	1620
accatacgtc	ccaacaaggc	agtcatcaac	cagggcacca	aggtcaggta	ccgaatgatc	1680
caccaatttc	aagaccagt	ttacaaccat	caacagtccac	cttggacgtg	ttccgtcggt	1740
actgttagaca	gacatttgg	aatcgat	ttgaaggaga	aagtgaacaa	tccggcgca	1800
taattcgtgc	tagtaacaaa	ttcattgaag	aatttgattt	gccgattttt	ggtgtgacag	1860
ttgttcgacc	ggggatgaca	gacgggtgagg	ttttggagaa	catcatgccc	gaagatgcac	1920
catatcatga	catttgcag	ttcatttga	ggctcacatc	agaaaagtgt	actttctcag	1980

gagaggggcc agaagttagt gatttgaacg aaaaatgggg aacaattgt tactatgaga 2040
aaaatttgc aattggcgag aaaaaatgtt cgagaggaaa ttccacgtg gatggcggat 2100
tcatttgctc tgagaatcgt tacagtctcg gacttgagcc aaatccaatt agagaaccag 2160
tggcgtttaa agttcgtaaa gcaatagtgg atggaattcg cttttccctac aaaaaaagacg 2220
ggagtgtttg gcttcaaaac cgcatgaagt acccggtatt tgtcacttct gggtatctcg 2280
acgagaatac aggaggccta aagaaggata aagtgcacaa agtttacgga tgtgcgtcta 2340
tcaaaacgtt tggcttcaac gtttccaaac aaatcatcag agacgcgtt ctttccaagc 2400
aaatggcaac aatgtacttg caaggaaaat tgactccgt gaattatatc tacgagaaga 2460
agactcagga agagctgcga agggaaagcaa cacgaccac tgattcattg gccaaagtact 2520
gttgtgtccc tgtctcggtc tgcaaaggat ttggagaagc atacccagaa cgcccgtaa 2580
ttcatgattt tccagttgg attgagttga aaatcaacat tgcctacgt ttcatggatt 2640
caatctgcca gtacataacc aactgcttcg agccgctagg aatgaaagat ttgcaaaat 2700
tggaaatcaa cgtcagtgtat gactaaatga taacttttt cactcaccct actagatact 2760
gatttagtct tattccaaat catccaacga tatcaaactt tttcccttga actttgcata 2820
ctatgttatac acaagttca agcagttca atacaaaacat aggatatgtt aacaactttt 2880
gataagaatc aagttaccaa ctgttcattg tgagtttga gctgtataga aggacaatgt 2940
atccccatacc tcaatcttta atagtcatca gtcactggtc ccgcaccaat ttttcgatt 3000
cgcatatgtc atatattgca ccgtggccct ttttattgtt acttttaata tattttcttc 3060
ccaaacttgcg aatatgattt atgaaccacc attttgagta ataaatgtat tttttgtgg 3119

<210> 54

<211> 103

<212> PRT

<213> *Caenorhabditis elegans*

<400> 54

Lys	Lys	Thr	Thr	Thr	Arg	Arg	Asn	Ala	Trp	Gly	Asn	Met	Ser	Tyr	Ala
1					5					10				15	
Glu	Leu	Ile	Thr	Thr	Ala	Ile	Met	Ala	Ser	Pro	Glu	Lys	Arg	Leu	Thr
							20			25				30	
Leu	Ala	Gln	Val	Tyr	Glu	Trp	Met	Val	Gln	Asn	Val	Pro	Tyr	Phe	Arg
							35			40			45		
Asp	Lys	Gly	Asp	Ser	Asn	Ser	Ser	Ala	Gly	Trp	Lys	Asn	Ser	Ile	Arg
							50			55			60		
His	Asn	Leu	Ser	Leu	His	Ser	Arg	Phe	Met	Arg	Ile	Gln	Asn	Glu	Gly
							65			70			75		80
Ala	Gly	Lys	Ser	Ser	Trp	Trp	Val	Ile	Asn	Pro	Asp	Ala	Lys	Pro	Gly
							85			90				95	
Met	Asn	Pro	Arg	Arg	Thr	Arg									
					100										

<210> 55

<211> 41

<212> PRT

<213> *Caenorhabditis elegans*

<400> 55

Thr	Phe	Met	Asn	Thr	Pro	Asp	Asp	Val	Met	Met	Asn	Asp	Asp	Met	Glu
1					5				10					15	
Pro	Ile	Pro	Arg	Asp	Arg	Cys	Asn	Thr	Trp	Pro	Met	Arg	Arg	Pro	Gln
							20			25			30		
Leu	Glu	Pro	Pro	Leu	Asn	Ser	Ser	Pro							
							35			40					

<210> 56

<211> 109

<212> PRT

<213> Caenorhabditis elegans

<400> 56

Asp Asp Thr Val Ser Gly Lys Thr Thr Arg Arg Asn Ala Trp
1 5 10 15
Gly Asn Met Ser Tyr Ala Glu Leu Ile Thr Thr Ala Ile Met Ala Ser
20 25 30
Pro Glu Lys Arg Leu Thr Leu Ala Gln Val Tyr Glu Trp Met Val Gln
35 40 45
Asn Val Pro Tyr Phe Arg Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly
50 55 60
Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met
65 70 75 80
Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn
85 90 95
Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg
100 105

<210> 57

<211> 655

<212> PRT

<213> Homo sapiens

<400> 57

Met Ala Glu Ala Pro Gln Val Val Glu Ile Asp Pro Asp Phe Glu Pro
1 5 10 15
Leu Pro Arg Pro Arg Ser Cys Thr Trp Pro Leu Pro Arg Pro Glu Phe
20 25 30
Ser Gln Ser Asn Ser Ala Thr Ser Ser Pro Ala Pro Ser Gly Ser Ala
35 40 45
Ala Ala Asn Pro Asp Ala Ala Ala Gly Leu Pro Ser Ala Ser Ala Ala
50 55 60
Ala Val Ser Ala Asp Phe Met Ser Asn Leu Ser Leu Leu Glu Glu Ser
65 70 75 80
Glu Asp Phe Pro Gln Ala Pro Gly Ser Val Ala Ala Ala Val Ala Ala
85 90 95
Ala Ala Ala Ala Ala Ala Thr Gly Gly Leu Cys Gly Asp Phe Gln Gly
100 105 110
Pro Glu Ala Gly Cys Leu His Pro Ala Pro Pro Gln Pro Pro Pro Pro
115 120 125
Gly Pro Val Ser Gln His Pro Pro Val Pro Pro Ala Ala Ala Gly Pro
130 135 140
Leu Ala Gly Gln Pro Arg Lys Ser Ser Ser Ser Arg Arg Asn Ala Trp
145 150 155 160
Gly Asn Leu Ser Tyr Ala Asp Leu Ile Thr Lys Ala Ile Glu Ser Ser
165 170 175
Ala Glu Lys Arg Leu Thr Leu Ser Gln Ile Tyr Glu Trp Met Val Lys
180 185 190
Ser Val Pro Tyr Phe Lys Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly
195 200 205
Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Lys Phe Ile
210 215 220
Arg Val Gln Asn Glu Gly Thr Gly Lys Ser Ser Trp Trp Met Leu Asn
225 230 235 240
Pro Glu Gly Gly Lys Ser Gly Lys Ser Pro Arg Arg Arg Ala Ala Ser
245 250 255

ପ୍ରକାଶକ ପତ୍ର ମହିନେ ପରିଚୟ ୧୦

Met Asp Asn Asn Ser Lys Phe Ala Lys Ser Arg Ser Arg Ala Ala Lys
 260 265 270
 Lys Lys Ala Ser Leu Gln Ser Gly Gln Glu Gly Ala Gly Asp Ser Pro
 275 280 285
 Gly Ser Gln Phe Ser Lys Trp Pro Ala Ser Pro Gly Ser His Ser Asn
 290 295 300
 Asp Asp Phe Asp Asn Trp Ser Thr Phe Arg Pro Arg Thr Ser Ser Asn
 305 310 315 320
 Ala Ser Thr Ile Ser Gly Arg Leu Ser Pro Ile Met Thr Glu Gln Asp
 325 330 335
 Asp Leu Gly Glu Gly Asp Val His Ser Met Val Tyr Pro Pro Ser Ala
 340 345 350
 Ala Lys Met Ala Ser Thr Leu Pro Ser Leu Ser Glu Ile Ser Asn Pro
 355 360 365
 Glu Asn Met Glu Asn Leu Leu Asp Asn Leu Asn Leu Leu Ser Ser Pro
 370 375 380
 Thr Ser Leu Thr Val Ser Thr Gln Ser Ser Pro Gly Thr Met Met Gln
 385 390 395 400
 Gln Thr Pro Cys Tyr Ser Phe Ala Pro Pro Asn Thr Ser Leu Asn Ser
 405 410 415
 Pro Ser Pro Asn Tyr Gln Lys Tyr Thr Tyr Gly Gln Ser Ser Met Ser
 420 425 430
 Pro Leu Pro Gln Met Pro Ile Gln Thr Leu Gln Asp Asn Lys Ser Ser
 435 440 445
 Tyr Gly Gly Met Ser Gln Tyr Asn Cys Ala Pro Gly Leu Leu Lys Glu
 450 455 460
 Leu Leu Thr Ser Asp Ser Pro Pro His Asn Asp Ile Met Thr Pro Val
 465 470 475 480
 Asp Pro Gly Val Ala Gln Pro Asn Ser Arg Val Leu Gly Gln Asn Val
 485 490 495
 Met Met Gly Pro Asn Ser Val Met Ser Thr Tyr Gly Ser Gln Ala Ser
 500 505 510
 His Asn Lys Met Met Asn Pro Ser Ser His Thr His Pro Gly His Ala
 515 520 525
 Gln Gln Thr Ser Ala Val Asn Gly Arg Pro Leu Pro His Thr Val Ser
 530 535 540
 Thr Met Pro His Thr Ser Gly Met Asn Arg Leu Thr Gln Val Lys Thr
 545 550 555 560
 Pro Val Gln Val Pro Leu Pro His Pro Met Gln Met Ser Ala Leu Gly
 565 570 575
 Gly Tyr Ser Ser Val Ser Ser Cys Asn Gly Tyr Gly Arg Met Gly Leu
 580 585 590
 Leu His Gln Glu Lys Leu Pro Ser Asp Leu Asp Gly Met Phe Ile Glu
 595 600 605
 Arg Leu Asp Cys Asp Met Glu Ser Ile Ile Arg Asn Asp Leu Met Asp
 610 615 620
 Gly Asp Thr Leu Asp Phe Asn Phe Asp Asn Val Leu Pro Asn Gln Ser
 625 630 635 640
 The Pro His Ser Val Lys Thr Thr His Ser Trp Val Ser Gly
 645 650 655

<210> 58
<211> 98
<212> PRT
<213> *Caenorhabditis elegans*

<400> 58

Lys Pro Asn Pro Trp Gly Glu Ser Tyr Ser Asp Ile Ile Ala Lys
1 5 10 15
Ala Leu Glu Ser Ala Pro Asp Gly Arg Leu Lys Leu Asn Glu Ile Tyr
20 25 30
Gln Trp Phe Ser Asp Asn Ile Pro Tyr Phe Gly Glu Arg Ser Ser Pro
35 40 45
Glu Glu Ala Ala Gly Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu
50 55 60
His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser
65 70 75 80
Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg
85 90 95
Thr Arg

<210> 59
<211> 7
<212> PRT
<213> Caenorhabditis elegans

<400> 59
Trp Lys Asn Ser Ile Arg His
1 5

<210> 60
<211> 121
<212> PRT
<213> Caenorhabditis elegans

<400> 60
Gln Val Leu Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly
1 5 10 15
Val Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
20 25 30
Ser Lys Asp His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu
35 40 45
Arg Phe Pro Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly
50 55 60
Leu Leu Val Lys Asp Pro Thr Gln Arg Leu Gly Gly Pro Glu Asp
65 70 75 80
Ala Leu Glu Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu
85 90 95
Ala Thr Tyr Arg Lys Glu Ile Glu Pro Pro Tyr Lys Pro Asn Val Gln
100 105 110
Ser Glu Thr Asp Thr Ser Tyr Phe Asp
115 120

<210> 61
<211> 66
<212> PRT
<213> Caenorhabditis elegans

<400> 61
Thr Met Glu Asp Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe
1 5 10 15

DRAFT

Gly Lys Val Ile Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala
20 25 30
Ile Lys Ile Leu Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala
35 40 45
His Thr Leu Thr Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe
50 55 60
Leu Thr
65

<210> 62
<211> 45
<212> PRT
<213> *Caenorhabditis elegans*

<400> 62
Lys Leu Glu Asn Leu Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
1 5 10 15
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
20 25 30
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
35 40 45

<210> 63
<211> 57
<212> PRT
<213> *Caenorhabditis elegans*

<400> 63
Tyr Phe Gln Glu Leu Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys
1 5 10 15
Phe Val Met Gln Phe Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg
20 25 30
Lys Cys Gly Thr Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu
35 40 45
Ile Val Leu Ala Leu Gly Tyr Leu His
50 55

<210> 64
<211> 59
<212> PRT
<213> *Caenorhabditis elegans*

<400> 64
Ser Thr Phe Ala Ile Phe Tyr Phe Gln Thr Met Leu Phe Glu Lys Pro
1 5 10 15
Arg Pro Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile
20 25 30
Glu Arg Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile
35 40 45
His Ala Ile Glu Ser Ile Ser Lys Lys Tyr Lys
50 55

<210> 65
<211> 33

DEPARTMENT OF COMPUTER SCIENCE

<212> PRT
<213> *Caenorhabditis elegans*

<400> 65
Leu Gln Glu Leu Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe
1 5 10 15
Val Met Glu Phe Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg
20 25 30
Glu

<210> 66
<211> 21
<212> PRT
<213> *Caenorhabditis elegans*

<400> 66
Val Val Ile Glu Gly Trp Leu His Lys Lys Gly Glu His Ile Arg Asn
1 5 10 15
Trp Arg Pro Arg Phe
20

<210> 67
<211> 26
<212> PRT
<213> *Caenorhabditis elegans*

<400> 67
Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ser Glu Ile Val Leu Ala
1 5 10 15
Leu Gly Tyr Leu His Ala Asn Ser Ile Val
20 25

<210> 68
<211> 39
<212> PRT
<213> *Caenorhabditis elegans*

<400> 68
Ile Arg Val Ser Phe Cys Lys Gly Phe Gly Glu Thr Tyr Ser Arg Leu
1 5 10 15
Lys Val Val Asn Leu Pro Cys Trp Ile Glu Ile Ile Leu His Glu Pro
20 25 30
Ala Asp Glu Tyr Asp Thr Val
35

<210> 69
<211> 45
<212> PRT
<213> *Caenorhabditis elegans*

<400> 69
Ser Arg Asn Ser Lys Ser Ser Gln Ile Arg Asn Thr Val Gly Ala Gly
1 5 10 15

EQUA = EQUA + EQUA

Ile Gln Leu Ala Tyr Glu Asn Gly Glu Leu Trp Leu Thr Val Leu Thr
20 25 30
Asp Gln Ile Val Phe Val Gln Cys Pro Phe Leu Asn Gln
35 40 45

<210> 70
<211> 29
<212> PRT
<213> *Caenorhabditis elegans*

<400> 70
Asn Glu Met Leu Asp Pro Glu Pro Lys Tyr Pro Lys Glu Glu Lys Pro
1 5 10 15
Trp Cys Thr Ile Phe Tyr Tyr Glu Leu Thr Val Arg Val
20 25

<210> 71
<211> 29
<212> PRT
<213> *Caenorhabditis elegans*

<400> 71
Gln Leu Gly Lys Ala Phe Glu Ala Lys Val Pro Thr Ile Thr Ile Asp
1 5 10 15
Gly Ala Thr Gly Ala Ser Asp Glu Cys Arg Met Ser Leu
20 25

<210> 72
<211> 105
<212> PRT
<213> *Caenorhabditis elegans*

<400> 72
Ser Pro Asp Asp Gly Leu Leu Asp Ser Ser Glu Glu Ser Arg Arg Arg
1 5 10 15
Gln Lys Thr Cys Arg Val Cys Gly Asp His Ala Thr Gly Tyr Asn Phe
20 25 30
Asn Val Ile Thr Cys Glu Ser Cys Lys Ala Phe Phe Arg Arg Asn Ala
35 40 45
Leu Arg Pro Lys Glu Phe Lys Cys Pro Tyr Ser Glu Asp Cys Glu Ile
50 55 60
Asn Ser Val Ser Arg Arg Phe Cys Gln Lys Cys Arg Leu Arg Lys Cys
65 70 75 80
Phe Thr Val Gly Met Lys Lys Glu Trp Ile Leu Asn Glu Glu Gln Leu
85 90 95
Arg Arg Arg Lys Asn Ser Arg Leu Asn
100 105

<210> 73
<211> 89
<212> PRT
<213> *Caenorhabditis elegans*

<400> 73

Leu Asp Ser Ser Glu Glu Ser Arg Arg Gln Lys Thr Cys Arg Val
1 5 10 15
Cys Gly Asp His Ala Thr Gly Tyr Asn Phe Asn Val Ile Thr Cys Glu
20 25 30
Ser Cys Lys Ala Phe Phe Arg Arg Asn Ala Leu Arg Pro Lys Glu Phe
35 40 45
Lys Cys Pro Tyr Ser Glu Asp Cys Glu Ile Asn Ser Val Ser Arg Arg
50 55 60
Phe Cys Gln Lys Cys Arg Leu Arg Lys Cys Phe Thr Val Gly Met Lys
65 70 75 80
Lys Glu Trp Ile Leu Asn Glu Glu Gln
85

<210> 74
<211> 73
<212> PRT
<213> *Caenorhabditis elegans*

<400> 74
Asp Ile Met Asn Ile Met Asp Val Thr Met Arg Arg Phe Val Lys Val
1 5 10 15
Ala Lys Gly Val Pro Ala Phe Arg Glu Val Ser Gln Glu Gly Lys Phe
20 25 30
Ser Leu Leu Lys Gly Gly Met Ile Glu Met Leu Thr Val Arg Gly Val
35 40 45
Thr Arg Tyr Asp Ala Ser Thr Asn Ser Phe Lys Thr Pro Thr Ile Lys
50 55 60
Gly Gln Asn Val Ser Val Asn Val Asp
65 70

<210> 75
<211> 112
<212> PRT
<213> *Caenorhabditis elegans*

<400> 75
Ser Gly Ser Leu Val Asp Leu Met Ile Lys Asn Leu Thr Ala Tyr Thr
1 5 10 15
Gln Gly Leu Asn Glu Thr Val Lys Asn Arg Thr Ala Glu Leu Glu Lys
20 25 30
Glu Gln Glu Lys Gly Asp Gln Leu Leu Met Glu Leu Leu Pro Lys Ser
35 40 45
Val Ala Asn Asp Leu Lys Asn Gly Ile Ala Val Asp Pro Lys Val Tyr
50 55 60
Glu Asn Ala Thr Ile Leu Tyr Ser Asp Ile Val Gly Phe Thr Ser Leu
65 70 75 80
Cys Ser Gln Ser Gln Pro Met Glu Val Val Thr Leu Leu Ser Gly Met
85 90 95
Tyr Gln Arg Phe Asp Leu Ile Ile Ser Gln Gln Gly Gly Tyr Lys Val
100 105 110

<210> 76
<211> 107
<212> PRT
<213> *Caenorhabditis elegans*

<400> 76
Met Glu Thr Ile Gly Asp Ala Tyr Cys Val Ala Ala Gly Leu Pro Val
1 5 10 15
Val Met Glu Lys Asp His Val Lys Ser Ile Cys Met Ile Ala Leu Leu
20 25 30
Gln Arg Asp Cys Leu His His Phe Glu Ile Pro His Arg Pro Gly Thr
35 40 45
Phe Leu Asn Cys Arg Trp Gly Phe Asn Ser Gly Pro Val Phe Ala Gly
50 55 60
Val Ile Gly Gln Lys Ala Pro Arg Tyr Ala Cys Phe Gly Glu Ala Val
65 70 75 80
Ile Leu Ala Ser Lys Met Glu Ser Ser Gly Val Glu Asp Arg Ile Gln
85 90 95
Met Thr Leu Ala Ser Gln Gln Leu Leu Glu Glu
100 105

<210> 77
<211> 43
<212> PRT
<213> *Caenorhabditis elegans*

<400> 77
Asp Ile Leu Lys Gly Leu Glu Tyr Ile His Ala Ser Ala Ile Asp Phe
1 5 10 15
His Gly Asn Leu Thr Leu His Asn Cys Met Leu Asp Ser His Trp Ile
20 25 30
Val Lys Leu Ser Gly Phe Gly Val Asn Arg Leu
35 40

<210> 78
<211> 15
<212> PRT
<213> *Caenorhabditis elegans*

<400> 78
Asp Met Tyr Ser Phe Gly Val Ile Leu His Glu Ile Ile Leu Lys
1 5 10 15

<210> 79
<211> 67
<212> PRT
<213> *Caenorhabditis elegans*

<400> 79
Ala Ile Lys Ile Asn Val Asp Asp Pro Ala Ser Thr Glu Asn Leu Asn
1 5 10 15
Tyr Leu Met Glu Ala Asn Ile Met Lys Asn Phe Lys Thr Asn Phe Ile
20 25 30
Val Gln Leu Tyr Gly Val Ile Ser Thr Val Gln Pro Ala Met Val Val
35 40 45
Met Glu Met Met Asp Leu Gly Asn Leu Arg Asp Tyr Leu Arg Ser Lys
50 55 60
Arg Glu Asp
65

<210> 80
<211> 54
<212> PRT
<213> *Caenorhabditis elegans*

<400> 80
Val Ile Lys Lys Pro Glu Cys Cys Glu Asn Tyr Trp Tyr Lys Val Met
1 5 10 15
Lys Met Cys Trp Arg Tyr Ser Pro Arg Asp Arg Pro Thr Phe Leu Gln
20 25 30
Leu Val His Leu Leu Ala Ala Glu Ala Ser Pro Glu Phe Arg Asp Leu
35 40 45
Ser Phe Val Leu Thr Asp
50

<210> 81
<211> 69
<212> PRT
<213> *Caenorhabditis elegans*

<400> 81
Lys Gln Asp Ser Gly Met Ala Ser Glu Leu Lys Asp Ile Phe Ala Asn
1 5 10 15
Ile His Thr Ile Thr Gly Tyr Leu Leu Val Arg Gln Ser Ser Pro Phe
20 25 30
Ile Ser Leu Asn Met Phe Arg Asn Leu Arg Arg Ile Glu Ala Lys Ser
35 40 45
Leu Phe Arg Asn Leu Tyr Ala Ile Thr Val Phe Glu Asn Pro Asn Leu
50 55 60
Lys Lys Leu Phe Asp
65

<210> 82
<211> 52
<212> PRT
<213> *Caenorhabditis elegans*

<400> 82
Phe Pro His Leu Arg Glu Ile Thr Gly Thr Leu Leu Val Phe Glu Thr
1 5 10 15
Glu Gly Leu Val Asp Leu Arg Lys Ile Phe Pro Asn Leu Arg Val Ile
20 25 30
Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg Asn
35 40 45
Pro Asp Leu Glu
50

<210> 83
<211> 46
<212> PRT
<213> *Caenorhabditis elegans*

<400> 83
Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn Gly Gly Val Arg
1 5 10 15

Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr Ile Asp Trp Lys
20 25 30
His Leu Ile Thr Ser Ser Ile Asn Asp Val Val Val Asp Asn
35 40 45

<210> 84
<211> 36
<212> PRT
<213> Caenorhabditis elegans

<400> 84
Tyr Asn Ala Asp Asp Trp Glu Leu Arg Gln Asp Asp Val Val Leu Gly
1 5 10 15
Gln Gln Cys Gly Glu Gly Ser Phe Gly Lys Val Tyr Leu Gly Thr Gly
20 25 30
Asn Asn Val Val
35

<210> 85
<211> 24
<212> PRT
<213> Caenorhabditis elegans

<400> 85
Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly
1 5 10 15
Phe Gly Glu Ala Tyr Pro Glu Arg
20

<210> 86
<211> 13
<212> PRT
<213> Caenorhabditis elegans

<400> 86
Gly Trp Asp Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala
1 5 10

<210> 87
<211> 121
<212> PRT
<213> Homo sapiens

<400> 87
Glu Val Leu Glu Asp Asn Asp Tyr Gly Arg Ala Val Asp Trp Trp Gly
1 5 10 15
Leu Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
20 25 30
Asn Gln Asp His Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Glu Ile
35 40 45
Arg Phe Pro Arg Thr Leu Gly Pro Glu Ala Lys Ser Leu Leu Ser Gly
50 55 60
Leu Leu Lys Lys Asp Pro Thr Gln Arg Leu Gly Gly Ser Glu Asp
65 70 75 80

Ala Lys Glu Ile Met Gln His Arg Phe Phe Ala Asn Ile Val Trp Gln
85 90 95
Asp Val Tyr Glu Lys Lys Leu Ser Pro Pro Phe Lys Pro Gln Val Thr
100 105 110
Ser Glu Thr Asp Thr Arg Tyr Phe Asp
115 120

<210> 88
<211> 121
<212> PRT
<213> *Caenorhabditis elegans*

<400> 88
Gln Val Leu Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly
1 5 10 15
Val Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
20 25 30
Ser Lys Asp His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu
35 40 45
Arg Phe Pro Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly
50 55 60
Leu Leu Val Lys Asp Pro Thr Gln Arg Leu Gly Gly Pro Glu Asp
65 70 75 80
Ala Leu Glu Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu
85 90 95
Ala Thr Tyr Arg Lys Glu Ile Glu Pro Pro Tyr Lys Pro Asn Val Gln
100 105 110
Ser Glu Thr Asp Thr Ser Tyr Phe Asp
115 120

<210> 89
<211> 66
<212> PRT
<213> *Homo sapiens*

<400> 89
Thr Met Asn Glu Phe Glu Tyr Leu Lys Leu Leu Gly Lys Gly Thr Phe
1 5 10 15
Gly Lys Val Ile Leu Val Lys Glu Lys Ala Thr Gly Arg Tyr Tyr Ala
20 25 30
Met Lys Ile Leu Lys Lys Glu Val Ile Val Ala Lys Asp Glu Val Ala
35 40 45
His Thr Leu Thr Glu Asn Arg Val Leu Gln Asn Ser Arg His Pro Phe
50 55 60
Leu Thr
65

<210> 90
<211> 66
<212> PRT
<213> *Caenorhabditis elegans*

<400> 90
Thr Met Glu Asp Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe
1 5 10 15

Gly Lys Val Ile Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala
20 25 30
Ile Lys Ile Leu Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala
35 40 45
His Thr Leu Thr Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe
50 55 60
Leu Thr
65

<210> 91
<211> 45
<212> PRT
<213> Homo sapiens

<400> 91
Lys Leu Glu Asn Leu Met Leu Asp Lys Asp Gly His Ile Lys Ile Thr
1 5 10 15
Asp Phe Gly Leu Cys Lys Glu Gly Ile Lys Asp Gly Ala Thr Met Lys
20 25 30
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
35 40 45

<210> 92
<211> 45
<212> PRT
<213> Caenorhabditis elegans

<400> 92
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
1 5 10 15
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
20 25 30
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
35 40 45

<210> 93
<211> 57
<212> PRT
<213> Homo sapiens

<400> 93
Phe Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys
1 5 10 15
Phe Val Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser
20 25 30
Arg Glu Arg Val Phe Ser Glu Asp Arg Ala Arg Phe Tyr Gly Ala Glu
35 40 45
Ile Val Ser Ala Leu Asp Tyr Leu His
50 55

<210> 94
<211> 57
<212> PRT
<213> Caenorhabditis elegans

PDB ID: 1B2D

<400> 94
Tyr Phe Gln Glu Leu Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys
1 5 10 15
Phe Val Met Gln Phe Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg
20 25 30
Lys Cys Gly Thr Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu
35 40 45
Ile Val Leu Ala Leu Gly Tyr Leu His
50 55

<210> 95
<211> 59
<212> PRT
<213> Homo sapiens

<400> 95
Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys Thr Glu Arg Pro
1 5 10 15
Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp Thr Thr Val Ile
20 25 30
Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg Glu Glu Trp Ala
35 40 45
Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys
50 55

<210> 96
<211> 59
<212> PRT
<213> Caenorhabditis elegans

<400> 96
Ser Thr Phe Ala Ile Phe Tyr Phe Gln Thr Met Leu Phe Glu Lys Pro
1 5 10 15
Arg Pro Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile
20 25 30
Glu Arg Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile
35 40 45
His Ala Ile Glu Ser Ile Ser Lys Lys Tyr Lys
50 55

<210> 97
<211> 33
<212> PRT
<213> Homo sapiens

<400> 97
Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys Phe
1 5 10 15
Val Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser Arg
20 25 30
Glu

<210> 98

<211> 33
 <212> PRT
 <213> *Caenorhabditis elegans*

 <400> 98
 Leu Gln Glu Leu Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe
 1 5 10 15
 Val Met Glu Phe Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg
 20 25 30
 Glu

<210> 99
 <211> 473
 <212> PRT
 <213> *Homo sapiens*

 <400> 99
 Met Leu Gly Thr Val Lys Met Glu Gly His Glu Thr Ser Asp Trp Asn
 1 5 10 15
 Ser Tyr Tyr Ala Asp Thr Gln Glu Ala Tyr Ser Ser Val Pro Val Ser
 20 25 30
 Asn Met Asn Ser Gly Leu Gly Ser Met Asn Ser Met Asn Thr Tyr Met
 35 40 45
 Thr Met Asn Thr Met Thr Ser Gly Asn Met Thr Pro Ala Ser Phe
 50 55 60
 Asn Met Ser Tyr Ala Asn Pro Ala Leu Gly Ala Gly Leu Ser Pro Gly
 65 70 75 80
 Ala Val Ala Gly Met Pro Gly Gly Ser Ala Gly Ala Met Asn Ser Met
 85 90 95
 Thr Ala Ala Gly Val Thr Ala Met Gly Thr Ala Leu Ser Pro Ser Gly
 100 105 110
 Met Gly Ala Met Gly Ala Gln Gln Ala Ala Ser Met Met Asn Gly Leu
 115 120 125
 Gly Pro Tyr Ala Ala Ala Met Asn Pro Cys Met Ser Pro Met Ala Tyr
 130 135 140
 Ala Pro Ser Asn Leu Gly Arg Ser Arg Ala Gly Gly Gly Asp Ala
 145 150 155 160
 Lys Thr Phe Lys Arg Ser Tyr Pro His Ala Lys Pro Pro Tyr Ser Tyr
 165 170 175
 Ile Ser Leu Ile Thr Met Ala Ile Gln Arg Ala Pro Ser Lys Met Leu
 180 185 190
 Thr Leu Ser Glu Ile Tyr Gln Trp Ile Met Asp Leu Phe Pro Tyr Tyr
 195 200 205
 Arg Gln Asn Gln Gln Arg Trp Gln Asn Ser Ile Arg His Ser Leu Ser
 210 215 220
 Phe Asn Asp Cys Phe Val Lys Val Ala Arg Ser Pro Asp Lys Pro Gly
 225 230 235 240
 Lys Gly Ser Tyr Trp Thr Leu His Pro Asp Ser Gly Asn Met Phe Glu
 245 250 255
 Asn Gly Cys Tyr Leu Arg Arg Gln Lys Arg Phe Lys Cys Glu Lys Gln
 260 265 270
 Pro Gly Ala Gly Gly Gly Gly Ser Gly Ser Gly Gly Ser Gly Ala
 275 280 285
 Lys Gly Gly Pro Glu Ser Arg Lys Asp Pro Ser Gly Ala Ser Asn Pro
 290 295 300
 Ser Ala Asp Ser Pro Leu His Arg Gly Val His Gly Lys Thr Gly Gln

305	310	315	320
Leu Glu Gly Ala Pro Ala Pro Gly Pro Ala Ala Ser Pro Gln Thr Leu			
325	330	335	
Asp His Ser Gly Ala Thr Ala Thr Gly Gly Ala Ser Glu Leu Lys Thr			
340	345	350	
Pro Ala Ser Ser Thr Ala Pro Pro Ile Ser Ser Gly Pro Gly Ala Leu			
355	360	365	
Ala Ser Val Pro Ala Ser His Pro Ala His Gly Leu Ala Pro His Glu			
370	375	380	
Ser Gln Leu His Leu Lys Gly Asp Pro His Tyr Ser Phe Asn His Pro			
385	390	395	400
Phe Ser Ile Asn Asn Leu Met Ser Ser Glu Gln Gln His Lys Leu			
405	410	415	
Asp Phe Lys Ala Tyr Glu Gln Ala Leu Gln Tyr Ser Pro Tyr Gly Ser			
420	425	430	
Thr Leu Pro Ala Ser Leu Pro Leu Gly Ser Ala Ser Val Thr Thr Arg			
435	440	445	
Ser Pro Ile Glu Pro Ser Ala Leu Glu Pro Ala Tyr Tyr Gln Gly Val			
450	455	460	
Tyr Ser Arg Pro Val Leu Asn Thr Ser			
465	470		

<210> 100
<211> 347
<212> PRT
<213> Homo sapiens

<400> 100			
Met Leu Gly Ser Val Lys Met Glu Ala His Asp Leu Ala Glu Trp Ser			
1	5	10	15
Tyr Tyr Pro Glu Ala Gly Glu Val Tyr Ser Pro Val Thr Pro Val Pro			
20	25	30	
Thr Met Ala Pro Leu Asn Ser Tyr Met Thr Leu Asn Pro Leu Ser Ser			
35	40	45	
Pro Tyr Pro Gly Gly Leu Pro Ala Ser Pro Leu Pro Ser Gly Pro Leu			
50	55	60	
Ala Pro Pro Ala Pro Ala Ala Pro Leu Gly Pro Thr Phe Pro Gly Leu			
65	70	75	80
Gly Leu Ser Gly Gly Ser Ser Ser Gly Tyr Gly Ala Pro Gly Pro			
85	90	95	
Gly Leu Val His Gly Lys Glu Met Pro Lys Gly Tyr Arg Ala Pro Ala			
100	105	110	
His Ala Lys Pro Pro Tyr Ser Tyr Ile Ser Leu Ile Thr Met Ala Ile			
115	120	125	
Gln Gln Ala Pro Gly Lys Val Leu Thr Leu Ser Glu Ile Tyr Gln Trp			
130	135	140	
Ile Met Asp Leu Phe Pro Tyr Tyr Arg Asp Asn Gln Gln Arg Trp Gln			
145	150	155	160
Asn Ser Ile Arg His Ser Leu Ser Phe Asn Asp Cys Phe Val Lys Val			
165	170	175	
Ala Arg Ser Pro Asp Lys Pro Gly Lys Gly Ser Tyr Trp Ala Leu His			
180	185	190	
Pro Ser Ser Gly Asn Met Phe Glu Asn Gly Cys Tyr Leu Arg Arg Gln			
195	200	205	
Lys Arg Phe Lys Leu Glu Glu Lys Val Lys Lys Gly Gly Ser Gly Ala			
210	215	220	
Ser Thr Thr Arg Asn Gly Thr Gly Ser Ala Ala Ser Thr Thr Thr Pro			

© 2007-2010

225	230	235	240
Ala Ala Thr Val Thr Ser Pro Pro Gln Pro Pro Pro Pro Ala Pro Glu			
245	250	255	
Pro Glu Ala Gln Gly Gly Glu Asp Val Gly Ala Leu Asp Cys Gly Ser			
260	265	270	
Pro Ala Ser Ser Thr Pro Tyr Phe Thr Gly Leu Glu Leu Pro Gly Asp			
275	280	285	
Leu Lys Leu Asp Ala Pro Tyr Asn Phe Asn His Pro Phe Ser Ile Asn			
290	295	300	
Asn Leu Met Ser Glu Gln Thr Pro Ala Pro Pro Lys Leu Asp Val Gly			
305	310	315	320
Phe Gly Gly Tyr Gly Ala Glu Gly Gly Glu Pro Gly Val Tyr Tyr Gln			
325	330	335	
Gly Leu Tyr Ser Arg Ser Leu Leu Asn Ala Ser			
340	345		

<210> 101

<211> 635

<212> PRT

<213> Caenorhabditis elegans

<400> 101

Met Met Glu Met Leu Val Asp Gln Gly Thr Asp Ala Ser Ser Ser Ala			
1	5	10	15
Ser Thr Ser Thr Ser Ser Val Ser Arg Phe Gly Ala Asp Thr Phe Met			
20	25	30	
Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu Pro Ile Pro			
35	40	45	
Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Gln Leu Glu Pro			
50	55	60	
Pro Leu Asn Ser Ser Pro Ile Ile His Glu Gln Ile Pro Glu Glu Asp			
65	70	75	80
Ala Asp Leu Tyr Gly Ser Asn Glu Gln Cys Gly Gln Leu Gly Gly Ala			
85	90	95	
Ser Ser Asn Gly Ser Thr Ala Met Leu His Thr Pro Asp Gly Ser Asn			
100	105	110	
Ser His Gln Thr Ser Phe Pro Ser Glu Cys Tyr Thr Trp Pro Met Gln			
115	120	125	
Gln Tyr Ile Tyr Gln Glu Ser Ser Ala Thr Ile Pro His His His Leu			
130	135	140	
Asn Gln His Asn Asn Pro Tyr His Pro Met His Pro His His Gln Leu			
145	150	155	160
Pro His Met Gln Gln Leu Pro Gln Pro Leu Leu Asn Leu Asn Met Thr			
165	170	175	
Thr Leu Thr Ser Ser Gly Ser Ser Val Ala Ser Ser Ile Gly Gly Gly			
180	185	190	
Ala Gln Cys Ser Pro Cys Ala Ser Gly Ser Ser Thr Ala Ala Thr Asn			
195	200	205	
Ser Ser Gln Gln Gln Gln Thr Val Gly Gln Met Leu Ala Ala Ser Val			
210	215	220	
Pro Cys Ser Ser Ser Gly Met Thr Leu Gly Met Ser Leu Asn Leu Ser			
225	230	235	240
Gln Gly Gly Gly Pro Met Pro Ala Lys Lys Lys Arg Cys Arg Lys Lys			
245	250	255	
Pro Thr Asp Gln Leu Ala Gln Lys Lys Pro Asn Pro Trp Gly Glu Glu			
260	265	270	
Ser Tyr Ser Asp Ile Ile Ala Lys Ala Leu Glu Ser Ala Pro Asp Gly			

□□□□□□□□□□□□□□□

275	280	285
Arg Leu Lys Leu Asn Glu Ile Tyr Gln Trp Phe Ser Asp Asn Ile Pro		
290	295	300
Tyr Phe Gly Glu Arg Ser Ser Pro Glu Glu Ala Ala Gly Trp Lys Asn		
305	310	315
Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met Arg Ile Gln		
325	330	335
Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro Asp Ala		
340	345	350
Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser Asn Thr Ile		
355	360	365
Glu Thr Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg Gly Ala Lys		
370	375	380
Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His Ser Thr Leu		
385	390	395
Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser His Asp Leu		
405	410	415
Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val Pro Ser Ser		
420	425	430
Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly Ser Ser Ser		
435	440	445
Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp Leu Glu Phe		
450	455	460
Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser Asp Ile Val		
465	470	475
Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His Ile Gly Gly		
485	490	495
Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu Pro Ile Ala		
500	505	510
Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly Ser Cys Ala		
515	520	525
Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr Asn Phe Lys		
530	535	540
Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn Gly Gly Ile		
545	550	555
Thr Pro Ile Asn Trp Leu Ser Thr Ser Asn Ser Ser Pro Leu Pro Gly		
565	570	575
Ile Gln Ser Cys Gly Ile Val Ala Ala Gln His Thr Val Ala Ser Ser		
580	585	590
Ser Ala Leu Pro Ile Asp Leu Glu Asn Leu Thr Leu Pro Asp Gln Pro		
595	600	605
Leu Met Asp Thr Met Asp Val Asp Ala Leu Ile Arg His Glu Leu Ser		
610	615	620
Gln Ala Gly Gly Gln His Ile His Phe Asp Leu		
625	630	635

<210> 102

<211> 501

<212> PRT

<213> Homo sapiens

<400> 102

Met Arg Ile Gln Pro Gln Lys Ala Ala Ala Ile Ile Asp Leu Asp Pro		
1	5	10
Asp Phe Glu Pro Gln Ser Arg Pro Arg Ser Cys Thr Trp Pro Leu Pro		
20	25	30
Arg Pro Glu Ile Ala Asn Gln Pro Ser Glu Pro Pro Glu Val Glu Pro		

35	40	45
Asp Leu Gly Glu Lys Val His Thr Glu Gly Arg Ser Glu Pro Ile Leu		
50	55	60
Leu Pro Ser Arg Leu Ser Glu Pro Ala Gly Gly Pro Gln Pro Gly Ile		
65	70	75
Leu Gly Ala Val Thr Gly Pro Arg Lys Gly Gly Ser Arg Arg Asn Ala		
85	90	95
Trp Gly Asn Gln Ser Tyr Ala Glu Phe Ile Ser Gln Ala Ile Glu Ser		
100	105	110
Ala Pro Glu Lys Arg Leu Thr Leu Ala Gln Ile Tyr Glu Trp Met Val		
115	120	125
Arg Thr Val Pro Tyr Phe Lys Asp Lys Gly Asp Ser Asn Ser Ser Ala		
130	135	140
Gly Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Lys Phe		
145	150	155
Ile Lys Val His Asn Glu Ala Thr Gly Lys Ser Ser Trp Trp Met Leu		
165	170	175
Asn Pro Glu Gly Lys Ser Gly Lys Ala Pro Arg Arg Arg Ala Ala		
180	185	190
Ser Met Asp Ser Ser Ser Lys Leu Leu Arg Gly Arg Ser Lys Ala Pro		
195	200	205
Lys Lys Lys Pro Ser Val Leu Pro Ala Pro Pro Glu Gly Ala Thr Pro		
210	215	220
Thr Ser Pro Val Gly His Phe Ala Lys Trp Ser Gly Ser Pro Cys Ser		
225	230	235
Arg Asn Arg Glu Glu Ala Asp Met Trp Thr Thr Phe Arg Pro Arg Ser		
245	250	255
Ser Ser Asn Ala Ser Ser Val Ser Thr Arg Leu Ser Pro Leu Arg Pro		
260	265	270
Glu Ser Glu Val Leu Ala Glu Glu Ile Pro Ala Ser Val Ser Ser Tyr		
275	280	285
Ala Gly Gly Val Pro Pro Thr Leu Asn Glu Gly Leu Glu Leu Leu Asp		
290	295	300
Gly Leu Asn Leu Thr Ser Ser His Ser Leu Leu Ser Arg Ser Gly Leu		
305	310	315
Ser Gly Phe Ser Leu Gln His Pro Gly Val Thr Gly Pro Leu His Thr		
325	330	335
Tyr Ser Ser Ser Leu Phe Ser Pro Ala Glu Gly Pro Leu Ser Ala Gly		
340	345	350
Glu Gly Cys Phe Ser Ser Ser Gln Ala Leu Glu Ala Leu Leu Thr Ser		
355	360	365
Asp Thr Pro Pro Pro Ala Asp Val Leu Met Thr Gln Val Asp Pro		
370	375	380
Ile Leu Ser Gln Ala Pro Thr Leu Leu Leu Leu Gly Gly Leu Pro Ser		
385	390	395
Ser Ser Lys Leu Ala Thr Gly Val Gly Leu Cys Pro Lys Pro Leu Glu		
405	410	415
Ala Arg Gly Pro Ser Ser Leu Val Pro Thr Leu Ser Met Ile Ala Pro		
420	425	430
Pro Pro Val Met Ala Ser Ala Pro Ile Pro Lys Ala Leu Gly Thr Pro		
435	440	445
Val Leu Thr Pro Pro Thr Glu Ala Ala Ser Gln Asp Arg Met Pro Gln		
450	455	460
Asp Leu Asp Leu Asp Met Tyr Met Glu Asn Leu Glu Cys Asp Met Asp		
465	470	475
Asn Ile Ile Ser Asp Leu Met Asp Glu Gly Glu Gly Leu Asp Phe Asn		
485	490	495
Phe Glu Pro Asp Pro		

<210> 103
<211> 366
<212> PRT
<213> Homo sapiens

<400> 103
Arg Gly Ala Ile Arg Ile Glu Lys Asn Ala Asp Leu Cys Tyr Leu Ser
1 5 10 15
Thr Val Asp Trp Ser Leu Ile Leu Asp Ala Val Ser Asn Asn Tyr Ile
20 25 30
Val Gly Asn Lys Pro Pro Lys Glu Cys Gly Asp Leu Cys Pro Gly Thr
35 40 45
Met Glu Glu Lys Pro Met Cys Glu Lys Thr Thr Ile Asn Asn Glu Tyr
50 55 60
Asn Tyr Arg Cys Trp Thr Thr Asn Arg Cys Gln Lys Met Cys Pro Ser
65 70 75 80
Thr Cys Gly Lys Arg Ala Cys Thr Glu Asn Asn Glu Cys Cys His Pro
85 90 95
Glu Cys Leu Gly Ser Cys Ser Ala Pro Asp Asn Asp Thr Ala Cys Val
100 105 110
Ala Cys Arg His Tyr Tyr Ala Gly Val Cys Val Pro Ala Cys Pro
115 120 125
Pro Asn Thr Tyr Arg Phe Glu Gly Trp Arg Cys Val Asp Arg Asp Phe
130 135 140
Cys Ala Asn Ile Leu Ser Ala Glu Ser Ser Asp Ser Glu Gly Phe Val
145 150 155 160
Ile His Asp Gly Glu Cys Met Gln Glu Cys Pro Ser Gly Phe Ile Arg
165 170 175
Asn Gly Ser Gln Ser Met Tyr Cys Ile Pro Cys Glu Gly Pro Cys Pro
180 185 190
Lys Val Cys Glu Glu Lys Lys Thr Lys Thr Ile Asp Ser Val Thr
195 200 205
Ser Ala Gln Met Leu Gln Gly Cys Thr Ile Phe Lys Gly Asn Leu Leu
210 215 220
Ile Asn Ile Arg Arg Gly Asn Asn Ile Ala Ser Glu Leu Glu Asn Phe
225 230 235 240
Met Gly Leu Ile Glu Val Val Thr Gly Tyr Val Lys Ile Arg His Ser
245 250 255
His Ala Leu Val Ser Leu Ser Phe Leu Lys Asn Leu Arg Leu Ile Leu
260 265 270
Gly Glu Glu Gln Leu Glu Gly Asn Tyr Ser Phe Tyr Val Leu Asp Asn
275 280 285
Gln Asn Leu Gln Gln Leu Trp Asp Trp Asp His Arg Asn Leu Thr Ile
290 295 300
Lys Ala Gly Lys Met Tyr Phe Ala Phe Asn Pro Lys Leu Cys Val Ser
305 310 315 320
Glu Ile Tyr Arg Met Glu Glu Val Thr Gly Thr Lys Gly Arg Gln Ser
325 330 335
Lys Gly Asp Ile Asn Thr Arg Asn Asn Gly Glu Arg Ala Ser Cys Glu
340 345 350
Ser Asp Val Leu His Phe Thr Ser Thr Thr Ser Lys Asn
355 360 365

<210> 104

<211> 370
<212> PRT
<213> Homo sapiens

<400> 104

Arg Gly Ser Val Arg Ile Glu Lys Asn Asn Glu Leu Cys Tyr Leu Ala
1 5 10 15
Thr Ile Asp Trp Ser Arg Ile Leu Asp Ser Val Glu Asp Asn Tyr Ile
20 25 30
Val Leu Asn Lys Asp Asp Asn Glu Glu Cys Gly Asp Ile Cys Pro Gly
35 40 45
Thr Ala Lys Gly Lys Thr Asn Cys Pro Ala Thr Val Ile Asn Gly Gln
50 55 60
Phe Val Glu Arg Cys Trp Thr His Ser His Cys Gln Lys Val Cys Pro
65 70 75 80
Thr Ile Cys Lys Ser His Gly Cys Thr Ala Glu Gly Leu Cys Cys His
85 90 95
Ser Glu Cys Leu Gly Asn Cys Ser Gln Pro Asp Asp Pro Thr Lys Cys
100 105 110
Val Ala Cys Arg Asn Phe Tyr Leu Asp Gly Arg Cys Val Glu Thr Cys
115 120 125
Pro Pro Pro Tyr Tyr His Phe Gln Asp Trp Arg Cys Val Asn Phe Ser
130 135 140
Phe Cys Gln Asp Leu His His Lys Cys Lys Asn Ser Arg Arg Gln Gly
145 150 155 160
Cys His Gln Tyr Val Ile His Asn Asn Lys Cys Ile Pro Glu Cys Pro
165 170 175
Ser Gly Tyr Thr Met Asn Ser Ser Asn Leu Leu Cys Thr Pro Cys Leu
180 185 190
Gly Pro Cys Pro Lys Val Cys His Leu Leu Glu Gly Glu Lys Thr Ile
195 200 205
Asp Ser Val Thr Ser Ala Gln Glu Leu Arg Gly Cys Thr Val Ile Asn
210 215 220
Gly Ser Leu Ile Ile Asn Ile Arg Gly Gly Asn Asn Leu Ala Ala Glu
225 230 235 240
Leu Glu Ala Asn Leu Gly Leu Ile Glu Glu Ile Ser Gly Tyr Leu Lys
245 250 255
Ile Arg Arg Ser Tyr Ala Leu Val Ser Leu Ser Phe Phe Arg Lys Leu
260 265 270
Arg Leu Ile Arg Gly Glu Thr Leu Glu Ile Gly Asn Tyr Ser Phe Tyr
275 280 285
Ala Leu Asp Asn Gln Asn Leu Arg Gln Leu Trp Asp Trp Ser Lys His
290 295 300
Asn Leu Thr Ile Thr Gln Gly Lys Leu Phe Phe His Tyr Asn Pro Lys
305 310 315 320
Leu Cys Leu Ser Glu Ile His Lys Met Glu Glu Val Ser Gly Thr Lys
325 330 335
Gly Arg Gln Glu Arg Asn Asp Ile Ala Leu Lys Thr Asn Gly Asp Gln
340 345 350
Ala Ser Cys Glu Asn Glu Leu Leu Lys Phe Ser Tyr Ile Arg Thr Ser
355 360 365
Phe Asp
370

<210> 105
<211> 383
<212> PRT

<213> Drosophila melanogaster

<400> 105
Arg Gly Gly Val Arg Ile Glu Lys Asn His Lys Leu Cys Tyr Asp Arg
1 5 10 15
Thr Ile Asp Trp Leu Glu Ile Leu Ala Glu Asn Glu Ser Gln Leu Val
20 25 30
Val Leu Thr Glu Asn Gly Lys Glu Lys Glu Cys Ser Leu Ser Lys Cys
35 40 45
Pro Gly Glu Ile Arg Ile Glu Glu Gly His Asp Asn Thr Ala Ile Glu
50 55 60
Gly Glu Leu Asn Ala Ser Cys Gln Leu His Asn Asn Arg Arg Leu Cys
65 70 75 80
Trp Asn Ser Lys Leu Cys Gln Thr Lys Cys Pro Glu Lys Cys Arg Asn
85 90 95
Asn Cys Ile Asp Glu His Thr Cys Cys Ser Gln Asp Cys Leu Gly Gly
100 105 110
Cys Val Ile Asp Lys Asn Gly Asn Glu Ser Cys Ile Ser Cys Arg Asn
115 120 125
Val Ser Phe Asn Asn Ile Cys Met Asp Ser Cys Pro Lys Gly Tyr Tyr
130 135 140
Gln Phe Asp Ser Arg Cys Val Thr Ala Asn Glu Cys Ile Thr Leu Thr
145 150 155 160
Lys Phe Glu Thr Asn Ser Val Tyr Ser Gly Ile Pro Tyr Asn Gly Gln
165 170 175
Cys Ile Thr His Cys Pro Thr Gly Gln Lys Ser Glu Asn Lys Arg
180 185 190
Met Cys Glu Pro Cys Pro Gly Gly Lys Cys Asp Lys Glu Cys Ser Ser
195 200 205
Gly Leu Ile Asp Ser Leu Glu Arg Ala Arg Glu Phe His Gly Cys Thr
210 215 220
Ile Ile Thr Gly Thr Glu Pro Leu Thr Ile Ser Ile Lys Arg Glu Ser
225 230 235 240
Gly Ala His Val Met Asp Glu Leu Lys Tyr Gly Leu Ala Ala Val His
245 250 255
Lys Ile Gln Ser Ser Leu Met Val His Leu Thr Tyr Gly Leu Lys Ser
260 265 270
Leu Lys Phe Phe Gln Ser Leu Thr Glu Ile Ser Gly Asp Pro Pro Met
275 280 285
Asp Ala Asp Lys Tyr Ala Leu Tyr Val Leu Asp Asn Arg Asp Leu Asp
290 295 300
Glu Leu Trp Gly Pro Asn Gln Thr Val Phe Ile Arg Lys Gly Gly Val
305 310 315 320
Phe Phe His Phe Asn Pro Lys Leu Cys Val Ser Thr Ile Asn Gln Leu
325 330 335
Leu Pro Met Leu Ala Ser Lys Pro Lys Phe Phe Glu Lys Ser Asp Glu
340 345 350
Gly Ala Asp Ser Asn Gly Asn Arg Gly Ser Cys Gly Thr Ala Val Leu
355 360 365
Asn Val Thr Leu Gln Ser Val Gly Ala Asn Ser Ala Ser Leu Asn
370 375 380

<210> 106

<211> 381

<212> PRT

<213> Caenorhabditis elegans

Homo sapiens

<400> 106

Asn Gly Gly Val Arg Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys
1 5 10 15
Thr Ile Asp Trp Lys His Leu Ile Thr Ser Ser Ile Asn Asp Val Val
20 25 30
Val Asp Asn Ala Ala Glu Tyr Ala Val Thr Glu Thr Gly Leu Met Cys
35 40 45
Pro Arg Gly Ala Cys Glu Glu Asp Lys Gly Glu Ser Lys Cys His Tyr
50 55 60
Leu Glu Glu Lys Asn Gln Glu Gln Gly Val Glu Arg Val Gln Ser Cys
65 70 75 80
Trp Ser Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu
85 90 95
Pro Thr Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys
100 105 110
His Asp Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala
115 120 125
Cys His Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys
130 135 140
Cys Asp Ala His Leu Tyr Leu Leu Gln Arg Arg Cys Val Thr Arg
145 150 155 160
Glu Gln Cys Leu Gln Leu Asn Pro Val Leu Ser Asn Lys Thr Val Pro
165 170 175
Ile Lys Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr
180 185 190
Gln Ile Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys
195 200 205
Cys Glu Ile Val Cys Glu Ile Asn His Val Ile Asp Thr Phe Pro Lys
210 215 220
Ala Gln Ala Ile Arg Leu Cys Asn Ile Ile Asp Gly Asn Leu Thr Ile
225 230 235 240
Glu Ile Arg Gly Lys Gln Asp Ser Gly Met Ala Ser Glu Leu Lys Asp
245 250 255
Ile Phe Ala Asn Ile His Thr Ile Thr Gly Tyr Leu Leu Val Arg Gln
260 265 270
Ser Ser Pro Phe Ile Ser Leu Asn Met Phe Arg Asn Leu Arg Arg Ile
275 280 285
Glu Ala Lys Ser Leu Phe Arg Asn Leu Tyr Ala Ile Thr Val Phe Glu
290 295 300
Asn Pro Asn Leu Lys Lys Leu Phe Asp Ser Thr Thr Asp Leu Thr Leu
305 310 315 320
Asp Arg Gly Thr Val Ser Ile Ala Asn Asn Lys Met Leu Cys Phe Lys
325 330 335
Tyr Ile Lys Gln Leu Met Ser Lys Leu Asn Ile Pro Leu Asp Pro Ile
340 345 350
Asp Gln Ser Glu Gly Thr Asn Gly Glu Lys Ala Ile Cys Glu Asp Met
355 360 365
Ala Ile Asn Val Ser Ile Thr Ala Val Asn Ala Asp Ser
370 375 380

<210> 107

<211> 370

<212> PRT

<213> Homo sapiens

<400> 107

Ala Leu Pro Val Ala Val Leu Ile Val Gly Gly Leu Val Ile Met

© 1996-2000, BioEdit Software

1	5	10	15
Leu Tyr Val Phe His Arg Lys Arg Asn Asn Ser Arg Leu Gly Asn Gly			
20	25	30	
Val Leu Tyr Ala Ser Val Asn Pro Glu Tyr Phe Ser Ala Ala Asp Val			
35	40	45	
Tyr Val Pro Asp Glu Trp Glu Val Ala Arg Glu Lys Ile Thr Met Ser			
50	55	60	
Arg Glu Leu Gly Gln Gly Ser Phe Gly Met Val Tyr Glu Gly Val Ala			
65	70	75	80
Lys Gly Val Val Lys Asp Glu Pro Glu Thr Arg Val Ala Ile Lys Thr			
85	90	95	
Val Asn Glu Ala Ala Ser Met Arg Glu Arg Ile Glu Phe Leu Asn Glu			
100	105	110	
Ala Ser Val Met Lys Glu Phe Asn Cys His His Val Val Arg Leu Leu			
115	120	125	
Gly Val Val Ser Gln Gly Gln Pro Thr Leu Val Ile Met Glu Leu Met			
130	135	140	
Thr Arg Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu Met			
145	150	155	160
Glu Asn Asn Pro Val Leu Ala Pro Pro Ser Leu Ser Lys Met Ile Gln			
165	170	175	
Met Ala Gly Glu Ile Ala Asp Gly Met Ala Tyr Leu Asn Ala Asn Lys			
180	185	190	
Phe Val His Arg Asp Leu Ala Ala Arg Asn Cys Met Val Ala Glu Asp			
195	200	205	
Phe Thr Val Lys Ile Gly Asp Phe Gly Met Thr Arg Asp Ile Tyr Glu			
210	215	220	
Thr Asp Tyr Tyr Arg Lys Gly Lys Gly Leu Leu Pro Val Arg Trp			
225	230	235	240
Met Ser Pro Glu Ser Leu Lys Asp Gly Val Phe Thr Thr Tyr Ser Asp			
245	250	255	
Val Trp Ser Phe Gly Val Val Leu Trp Glu Ile Ala Thr Leu Ala Glu			
260	265	270	
Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln Val Leu Arg Phe Val Met			
275	280	285	
Glu Gly Gly Leu Leu Asp Lys Pro Asp Asn Cys Pro Asp Met Leu Phe			
290	295	300	
Glu Leu Met Arg Met Cys Trp Gln Tyr Asn Pro Lys Met Arg Pro Ser			
305	310	315	320
Phe Leu Glu Ile Ile Ser Ser Ile Lys Glu Glu Met Glu Pro Gly Phe			
325	330	335	
Arg Glu Val Ser Phe Tyr Tyr Ser Glu Glu Asn Lys Leu Pro Glu Pro			
340	345	350	
Glu Glu Leu Asp Leu Glu Pro Glu Asn Met Glu Ser Val Pro Leu Asp			
355	360	365	
Pro Ser			
370			

<210> 108

<211> 374

<212> PRT

<213> Homo sapiens

<400> 108

Ile Gly Pro Leu Ile Phe Val Phe Leu Phe Ser Val Val Ile Gly Ser			
1	5	10	15
Ile Tyr Leu Phe Leu Arg Lys Arg Gln Pro Asp Gly Pro Leu Gly Pro			

DRAFT 10/20/98

	20	25	30												
Leu	Tyr	Ala	Ser	Ser	Asn	Pro	Glu	Tyr	Leu	Ser	Ala	Ser	Asp	Val	Phe
35							40					45			
Pro	Cys	Ser	Val	Tyr	Val	Pro	Asp	Glu	Trp	Glu	Val	Ser	Arg	Glu	Lys
50						55				60					
Ile	Thr	Leu	Leu	Arg	Glu	Leu	Gly	Gln	Gly	Ser	Phe	Gly	Met	Val	Tyr
65						70				75			80		
Glu	Gly	Asn	Ala	Arg	Asp	Ile	Ile	Lys	Gly	Glu	Ala	Glu	Thr	Arg	Val
						85			90			95			
Ala	Val	Lys	Thr	Val	Asn	Glu	Ser	Ala	Ser	Leu	Arg	Glu	Arg	Ile	Glu
						100			105			110			
Phe	Leu	Asn	Glu	Ala	Ser	Val	Met	Lys	Gly	Phe	Thr	Cys	His	His	Val
						115			120			125			
Val	Arg	Leu	Leu	Gly	Val	Val	Ser	Lys	Gly	Gln	Pro	Thr	Leu	Val	Val
						130			135			140			
Met	Glu	Leu	Met	Ala	His	Gly	Asp	Leu	Lys	Ser	Tyr	Leu	Arg	Ser	Leu
145						150				155			160		
Arg	Pro	Glu	Ala	Glu	Asn	Asn	Pro	Gly	Arg	Pro	Pro	Pro	Thr	Leu	Gln
						165			170			175			
Glu	Met	Ile	Gln	Met	Ala	Ala	Glu	Ile	Ala	Asp	Gly	Met	Ala	Tyr	Leu
						180			185			190			
Asn	Ala	Lys	Lys	Phe	Val	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Cys	Met
						195			200			205			
Val	Ala	His	Asp	Phe	Thr	Val	Lys	Ile	Gly	Asp	Phe	Gly	Met	Thr	Arg
						210			215			220			
Asp	Ile	Tyr	Glu	Thr	Asp	Tyr	Tyr	Arg	Lys	Gly	Gly	Lys	Gly	Leu	Leu
225						230				235			240		
Pro	Val	Arg	Trp	Met	Ala	Pro	Glu	Ser	Leu	Lys	Asp	Gly	Val	Phe	Thr
						245			250			255			
Thr	Ser	Ser	Asp	Met	Trp	Ser	Phe	Gly	Val	Val	Leu	Trp	Glu	Ile	Thr
						260			265			270			
Ser	Leu	Ala	Glu	Gln	Pro	Tyr	Gln	Gly	Leu	Ser	Asn	Glu	Gln	Val	Leu
						275			280			285			
Lys	Phe	Val	Met	Asp	Gly	Gly	Tyr	Leu	Asp	Gln	Pro	Asp	Asn	Cys	Pro
						290			295			300			
Glu	Arg	Val	Thr	Asp	Leu	Met	Arg	Met	Cys	Trp	Gln	Phe	Asn	Pro	Lys
305						310				315			320		
Met	Arg	Pro	Thr	Phe	Leu	Glu	Ile	Val	Asn	Leu	Leu	Lys	Asp	Asp	Leu
						325			330			335			
His	Pro	Ser	Phe	Pro	Glu	Val	Ser	Phe	Phe	His	Ser	Glu	Glu	Asn	Lys
						340			345			350			
Ala	Pro	Glu	Ser	Glu	Glu	Leu	Met	Glu	Phe	Glu	Asp	Met	Glu	Asn	
						355			360			365			
Val	Pro	Leu	Asp	Arg	Ser										
						370									

<210> 109

<211> 384

<212> PRT

<213> Drosophila melanogaster

<400> 109

Gly	Ile	Gly	Leu	Ala	Phe	Leu	Ile	Val	Ser	Leu	Phe	Gly	Tyr	Val	Cys
1					5				10			15			
Tyr	Leu	His	Lys	Arg	Lys	Val	Pro	Ser	Asn	Asp	Leu	His	Met	Asn	Thr
							20		25			30			
Glu	Val	Asn	Pro	Phe	Tyr	Ala	Ser	Met	Gln	Tyr	Ile	Pro	Asp	Asp	Trp

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100

35	40	45
Glu Val Leu Arg Glu Asn Ile Ile Gln Leu Ala Pro Leu Gly Gln Gly		
50	55	60
Ser Phe Gly Met Val Tyr Glu Gly Ile Leu Lys Ser Phe Pro Pro Asn		
65	70	75
Gly Val Asp Arg Glu Cys Ala Ile Lys Thr Val Asn Glu Asn Ala Thr		
85	90	95
Asp Arg Glu Arg Thr Asn Phe Leu Ser Glu Ala Ser Val Met Lys Glu		
100	105	110
Phe Asp Thr Tyr His Val Val Arg Leu Leu Gly Val Cys Ser Arg Gly		
115	120	125
Gln Pro Ala Leu Val Val Met Glu Leu Met Lys Lys Gly Asp Leu Lys		
130	135	140
Ser Tyr Leu Arg Ala His Arg Pro Glu Glu Arg Asp Glu Ala Met Met		
145	150	155
Thr Tyr Leu Asn Arg Ile Gly Val Thr Gly Asn Val Gln Pro Pro Thr		
165	170	175
Tyr Gly Arg Ile Tyr Gln Met Ala Ile Glu Ile Ala Asp Gly Met Ala		
180	185	190
Tyr Leu Ala Ala Lys Lys Phe Val His Arg Asp Leu Ala Ala Arg Asn		
195	200	205
Cys Met Val Ala Asp Asp Leu Thr Val Lys Ile Gly Asp Phe Gly Met		
210	215	220
Thr Arg Asp Ile Tyr Glu Thr Asp Tyr Tyr Arg Lys Gly Thr Lys Gly		
225	230	235
Leu Leu Pro Val Arg Trp Met Pro Pro Glu Ser Leu Arg Asp Gly Val		
245	250	255
Tyr Ser Ser Ala Ser Asp Val Phe Ser Phe Gly Val Val Leu Trp Glu		
260	265	270
Met Ala Thr Leu Ala Ala Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln		
275	280	285
Val Leu Arg Tyr Val Ile Asp Gly Gly Val Met Glu Arg Pro Glu Asn		
290	295	300
Cys Pro Asp Phe Leu His Lys Leu Met Gln Arg Cys Trp His His Arg		
305	310	315
Ser Ser Ala Arg Pro Ser Phe Leu Asp Ile Ile Ala Tyr Leu Glu Pro		
325	330	335
Gln Cys Pro Asn Ser Gln Phe Lys Glu Val Ser Phe Tyr His Ser Glu		
340	345	350
Ala Gly Leu Gln His Arg Glu Lys Glu Arg Lys Glu Arg Asn Gln Leu		
355	360	365
Asp Ala Phe Ala Ala Val Pro Leu Asp Gln Asp Leu Gln Asp Arg Glu		
370	375	380

<210> 110

<211> 380

<212> PRT

<213> Caenorhabditis elegans

<400> 110

Gly Met Leu Leu Val Phe Leu Ile Leu Met Ser Ile Ala Gly Cys Ile		
1	5	10
Ile Tyr Tyr Tyr Ile Gln Val Arg Tyr Gly Lys Lys Val Lys Ala Leu		
20	25	30
Ser Asp Phe Met Gln Leu Asn Pro Glu Tyr Cys Val Asp Asn Lys Tyr		
35	40	45
Asn Ala Asp Asp Trp Glu Leu Arg Gln Asp Asp Val Val Leu Gly Gln		

DRAFT

50	55	60
Gln Cys Gly Glu Gly Ser Phe Gly Lys Val Tyr Leu Gly Thr Gly Asn		
65	70	75
Asn Val Val Ser Leu Met Gly Asp Arg Phe Gly Pro Cys Ala Ile Lys		80
85	90	95
Ile Asn Val Asp Asp Pro Ala Ser Thr Glu Asn Leu Asn Tyr Leu Met		
100	105	110
Glu Ala Asn Ile Met Lys Asn Phe Lys Thr Asn Phe Ile Val Gln Leu		
115	120	125
Tyr Gly Val Ile Ser Thr Val Gln Pro Ala Met Val Val Met Glu Met		
130	135	140
Met Asp Leu Gly Asn Leu Arg Asp Tyr Leu Arg Ser Lys Arg Glu Asp		
145	150	155
Glu Val Phe Asn Glu Thr Asp Cys Asn Phe Phe Asp Ile Ile Pro Arg		160
165	170	175
Asp Lys Phe His Glu Trp Ala Ala Gln Ile Cys Asp Gly Met Ala Tyr		
180	185	190
Leu Glu Ser Leu Lys Phe Cys His Arg Asp Leu Ala Ala Arg Asn Cys		
195	200	205
Met Ile Asn Arg Asp Glu Thr Val Lys Ile Gly Asp Phe Gly Met Ala		
210	215	220
Arg Asp Leu Phe Tyr His Asp Tyr Tyr Lys Pro Ser Gly Lys Arg Met		
225	230	235
Met Pro Val Arg Trp Met Ser Pro Glu Ser Leu Lys Asp Gly Lys Phe		240
245	250	255
Asp Ser Lys Ser Asp Val Trp Ser Phe Gly Val Val Leu Tyr Glu Met		
260	265	270
Val Thr Leu Gly Ala Gln Pro Tyr Ile Gly Leu Ser Asn Asp Glu Val		
275	280	285
Leu Asn Tyr Ile Gly Met Ala Arg Lys Val Ile Lys Lys Pro Glu Cys		
290	295	300
Cys Glu Asn Tyr Trp Tyr Lys Val Met Lys Met Cys Trp Arg Tyr Ser		
305	310	315
Pro Arg Asp Arg Pro Thr Phe Leu Gln Leu Val His Leu Leu Ala Ala		320
325	330	335
Glu Ala Ser Pro Glu Phe Arg Asp Leu Ser Phe Val Leu Thr Asp Asn		
340	345	350
Gln Met Ile Leu Asp Asp Ser Glu Ala Leu Asp Leu Asp Asp Ile Asp		
355	360	365
Asp Thr Asp Met Asn Asp Gln Val Val Glu Val Ala		
370	375	380

<210> 111

<211> 103

<212> PRT

<213> *Caenorhabditis elegans*

<400> 111

Asn Ile Asp Arg Glu Phe Asp Gln Lys Ala Cys Glu Ser Leu Val Lys		
1	5	10
Lys Leu Lys Asp Lys Lys Asn Asp Leu Gln Asn Leu Ile Asp Val Val		15
20	25	30
Leu Ser Lys Gly Thr Lys Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr		
35	40	45
Leu Asp Gly Arg Leu Gln Val His Gly Arg Lys Gly Phe Pro His Val		
50	55	60
Val Tyr Gly Lys Leu Trp Arg Phe Asn Glu Met Thr Lys Asn Glu Thr		

65 70 75 80
Arg His Val Asp His Cys Lys His Ala Phe Glu Met Lys Ser Asp Met
 85 90 95
Val Cys Val Asn Pro Tyr His
 100

<210> 112
<211> 104
<212> PRT
<213> Homo sapiens

<400> 112
Gly Gly Glu Ser Glu Thr Phe Ala Lys Arg Ala Ile Glu Ser Leu Val
1 5 10 15
Lys Lys Leu Lys Glu Lys Lys Asp Glu Leu Asp Ser Leu Ile Thr Ala
20 25 30
Ile Thr Thr Asn Gly Ala His Pro Ser Lys Cys Val Thr Ile Gln Arg
35 40 45
Thr Leu Asp Gly Arg Leu Gln Val Ala Gly Arg Lys Gly Phe Pro His
50 55 60
Val Ile Tyr Ala Arg Leu Trp Arg Trp Pro Asp Leu His Lys Asn Glu
65 70 75 80
Leu Lys His Val Lys Tyr Cys Gln Tyr Ala Phe Asp Leu Lys Cys Asp
85 90 95
Ser Val Cys Val Asn Pro Tyr His
100

<210> 113
<211> 205
<212> PRT
<213> Caenorhabditis elegans

<400> 113
Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys Lys Cys Ser
1 5 10 15
Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser Glu Asn Arg
20 25 30
Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val Ala Phe
35 40 45
Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr Lys Lys
50 55 60
Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val Phe Val
65 70 75 80
Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys Asp Lys
85 90 95
Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe Gly Phe Asn
100 105 110
Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys Gln Met Ala
115 120 125
Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr Ile Tyr Glu
130 135 140
Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg Thr Thr Asp
145 150 155 160
Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly Phe
165 170 175
Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys Pro Val Trp

180 185 190
Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp
195 200 205

<210> 114
<211> 212
<212> PRT
<213> Homo sapiens

<400> 114
Ile Ala Tyr Phe Glu Met Asp Val Gln Val Gly Glu Thr Phe Lys Val
1 5 10 15
Pro Ser Ser Cys Pro Ile Val Thr Val Asp Gly Tyr Val Asp Pro Ser
20 25 30
Gly Gly Asp Arg Phe Cys Leu Gly Gln Leu Ser Asn Val His Arg Thr
35 40 45
Glu Ala Ile Glu Arg Ala Arg Leu His Ile Gly Lys Gly Val Gln Leu
50 55 60
Glu Cys Lys Gly Glu Gly Asp Val Trp Val Arg Cys Leu Ser Asp His
65 70 75 80
Ala Val Phe Val Gln Ser Tyr Tyr Leu Asp Arg Glu Ala Gly Arg Ala
85 90 95
Pro Gly Asp Ala Val His Lys Ile Tyr Pro Ser Ala Tyr Ile Lys Val
100 105 110
Phe Asp Leu Arg Gln Cys His Arg Gln Met Gln Gln Ala Ala Thr
115 120 125
Ala Gln Ala Ala Ala Ala Gln Ala Ala Ala Val Ala Gly Asn Ile
130 135 140
Pro Gly Pro Gly Ser Val Gly Gly Ile Ala Pro Ala Ile Ser Leu Ser
145 150 155 160
Ala Ala Ala Gly Ile Gly Val Asp Asp Leu Arg Arg Leu Cys Ile Leu
165 170 175
Arg Met Ser Phe Val Lys Gly Trp Gly Pro Asp Tyr Pro Arg Gln Ser
180 185 190
Ile Lys Glu Thr Pro Cys Trp Ile Glu Ile His Leu His Arg Ala Leu
195 200 205
Gln Leu Leu Asp
210